

The diagram illustrates a computer system 100. At the top, three computer terminals are shown, labeled 125A, 125B, and 125C. Each terminal consists of a monitor and a base unit with a keyboard. These terminals are connected to a central computer system 100. The central system is represented by a large rectangle containing a PROCESSOR 105, INTERNAL STORAGE 110, and MEMORY 115. The PROCESSOR 105 is connected to both the INTERNAL STORAGE 110 and the MEMORY 115. Below the central system, a DATA RETRIEVING DEVICE 118 and a DISPLAY 120 are connected to the system via lines.

FIGURE 1

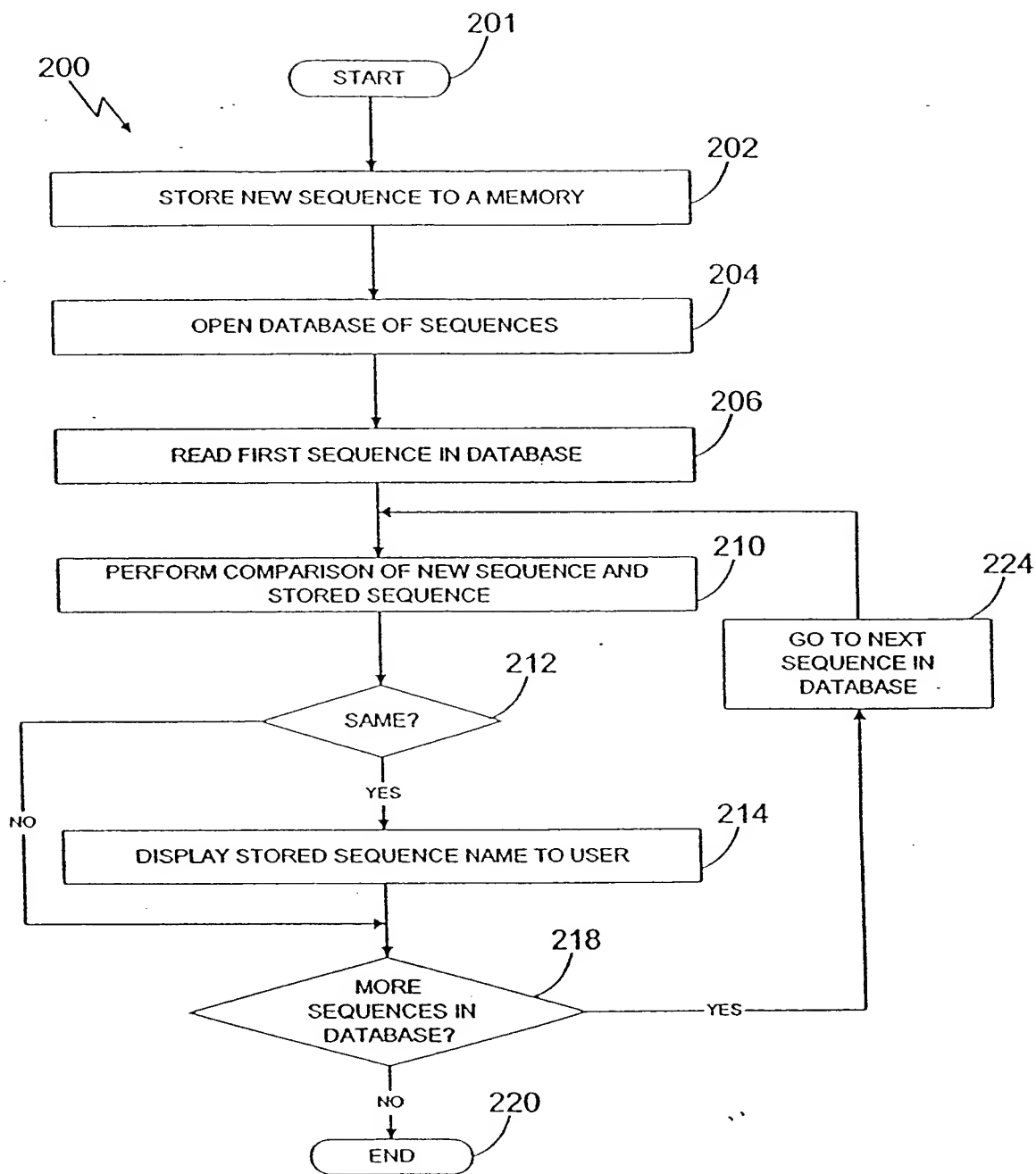


FIGURE 2

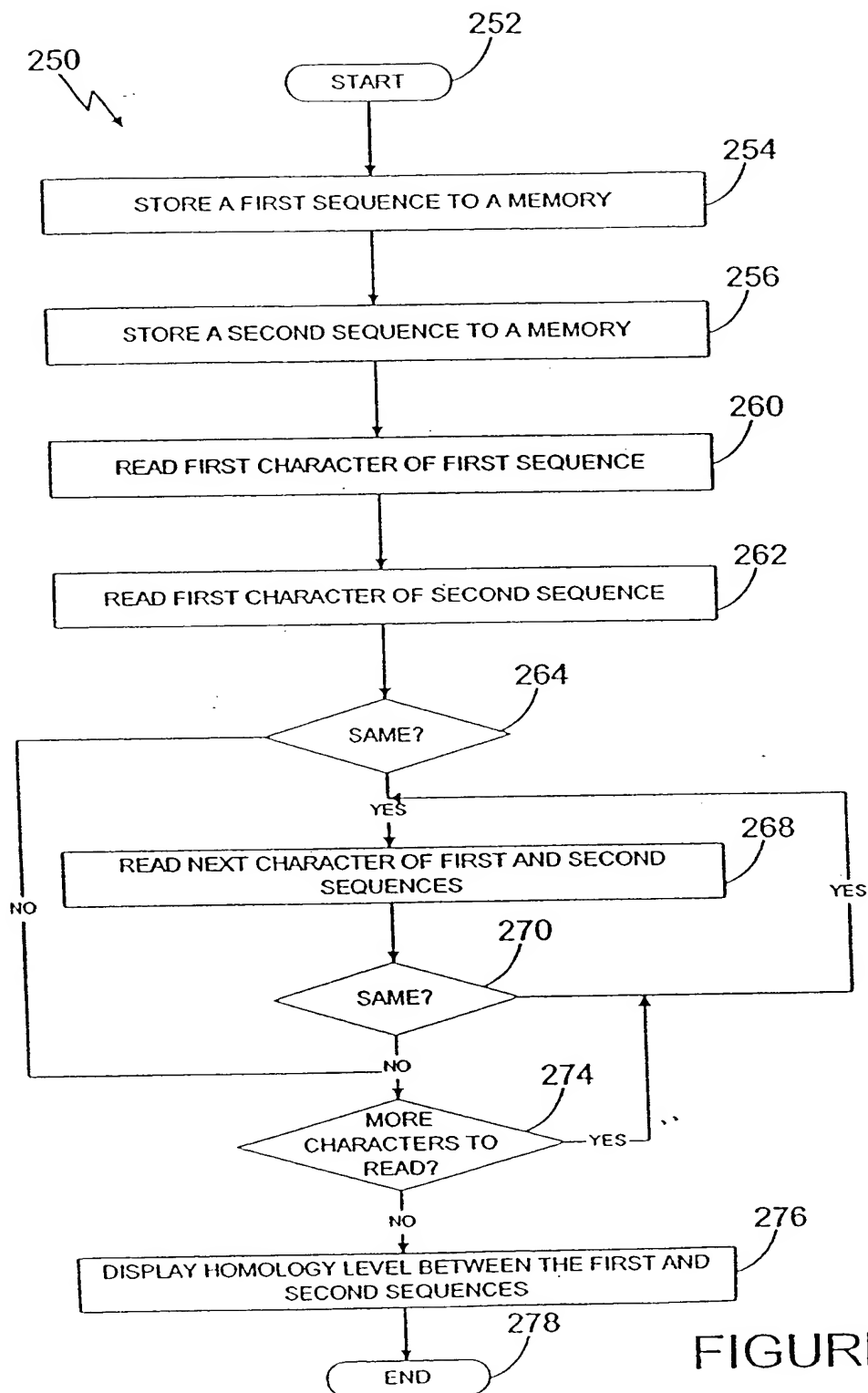


FIGURE 3

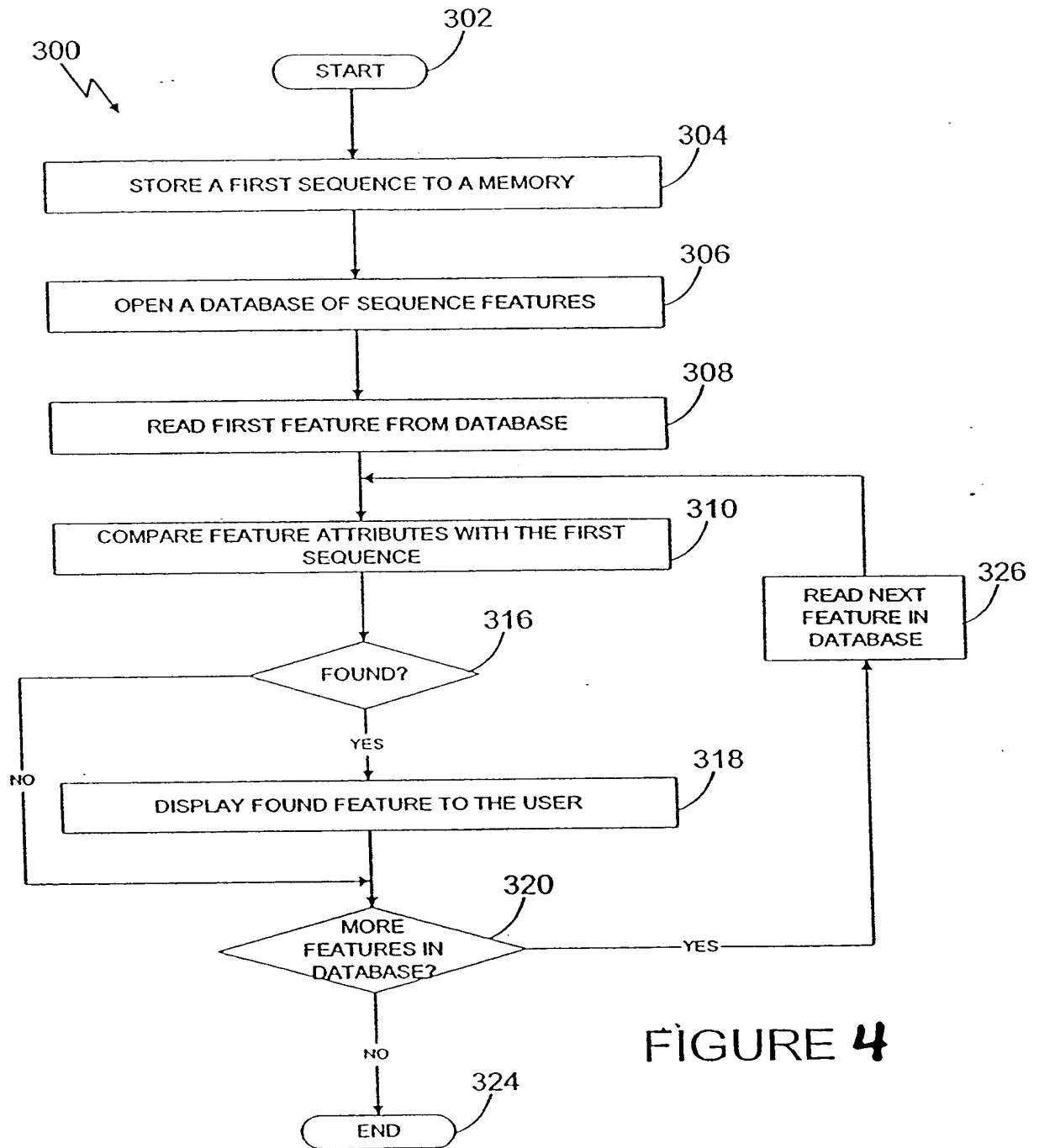
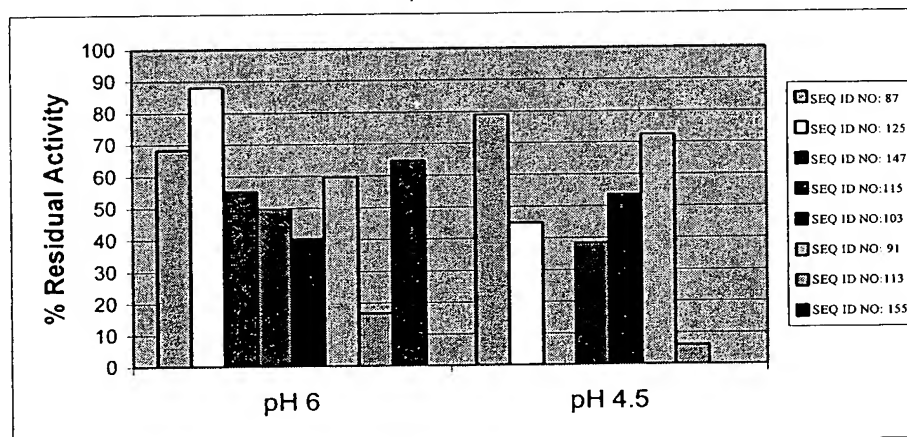
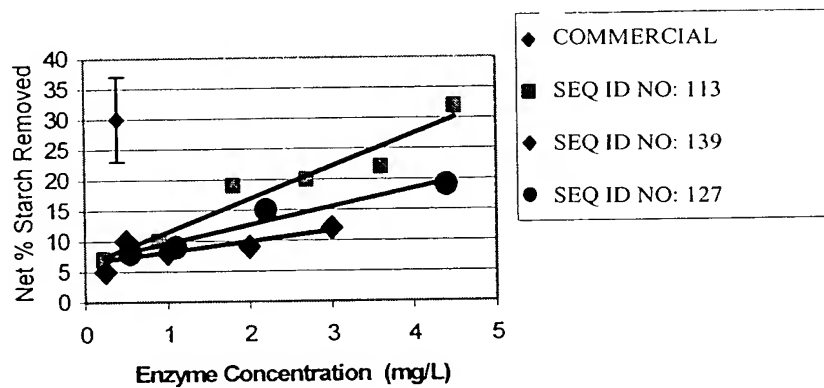


FIGURE 4



**Figure 5:** Residual activity of various amylases following heating to 90°C for 10 min.



**Figure 6.** Net percent starch removed vs. enzyme concentration in ADW wash test with bleach and chelators

**Figure 7:** Activity of parental amylases at pH 8, 40°C (black bars) in ADW formulation at 55°C (gray bars). Values are the average of 384 wells with error bars representing the standard deviation.

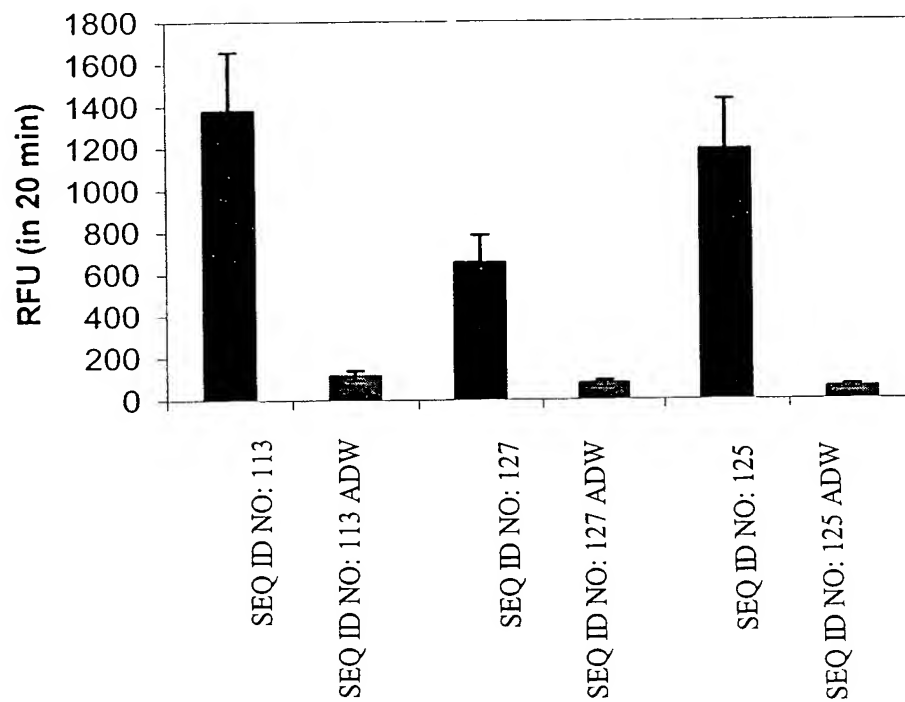
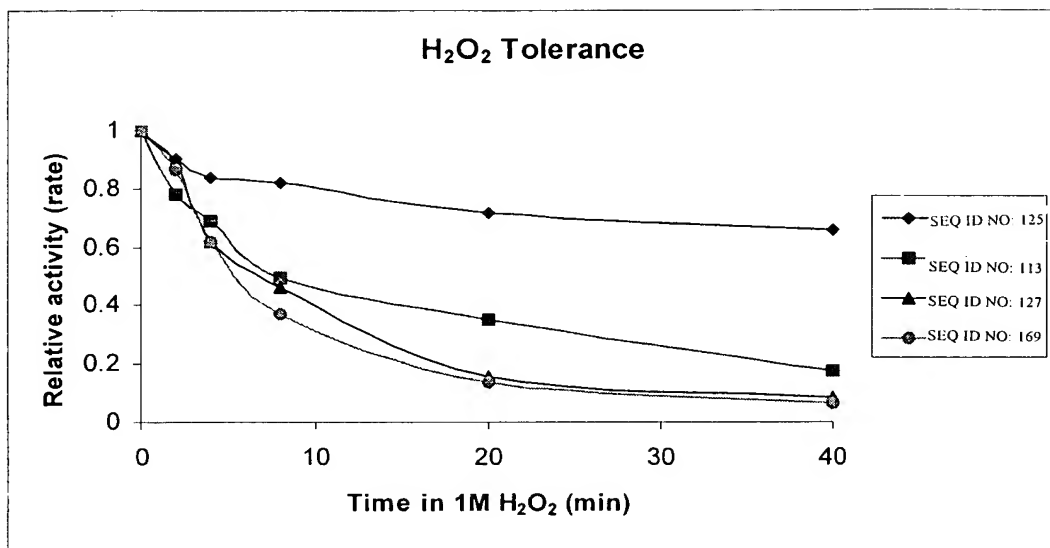


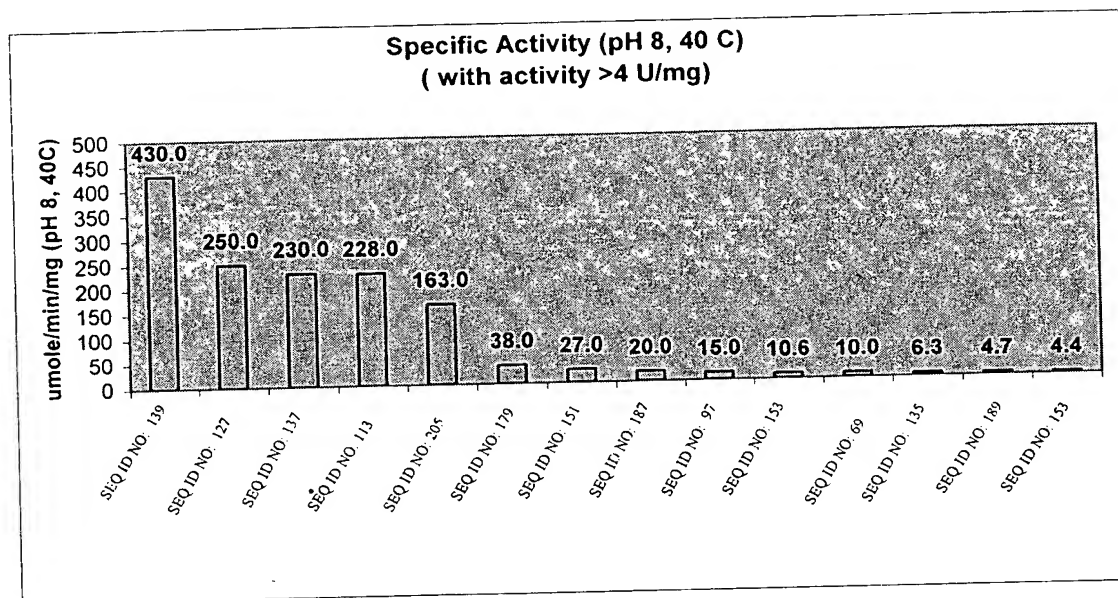
Figure 8:



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**Figure 9:** A graph of pH and temperature data for a selection of the amylases characterized: a) pH 8 and 40°C b) pH 10 and 50°C.

a)



b)

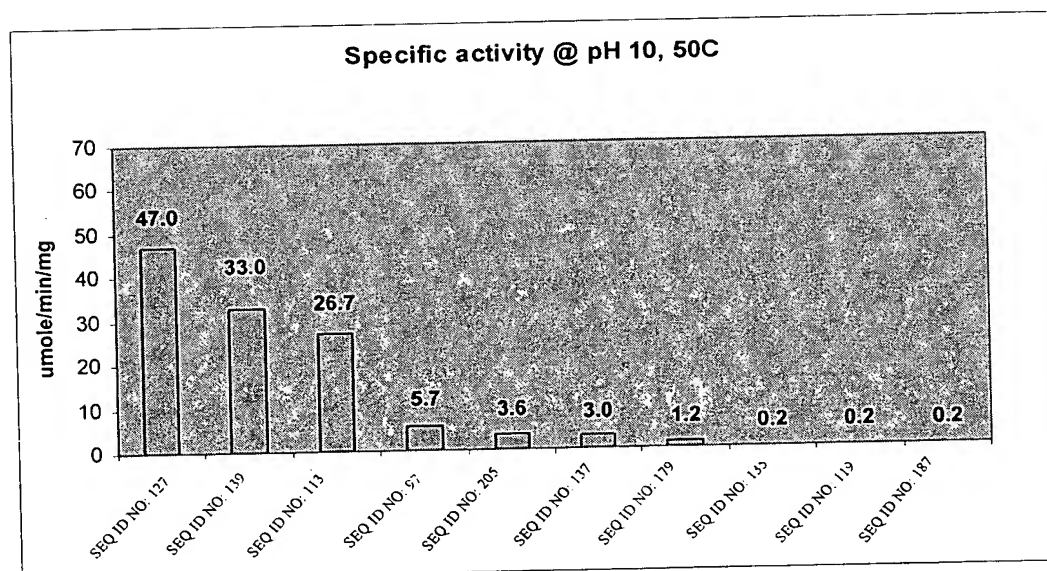


Figure 10: Alignments of the genes proposed to be used in reassembly

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1
SEQ ID NO.:113 (1) ----AANLNGTLMQYFEWYMPNDGQHWRKLNQDSAYLAEHGITAVWIPPAYKGTGTS-QADVGYGAYDLYDLGEFFHQKGTVR
SEQ ID NO.:127 (1) -QANTAPVNGTMMQYFEWDLPLNDGTLTKVKNEASSLSLGITALLWLPAYKGTGTS-QGDVGYGVYDLYDLGEFFHQKGTVR
SEQ ID NO.:115 (1) AKYSELEQGGVIMQAFYWDVPEGGIWDWTIRQKIPEWYDAGISAIWTPPASKMGGAYSMDYDPYDYFDLGEFFHQKGTVE

81
SEQ ID NO.:113 (76) TKYGTGKELQSAIKSLHSRDIINVYGDVVLNHKGGADATEDVTAVEVDPADRNRVISGEHRIKAWTHFHFPGRGSTYSDFK
SEQ ID NO.:127 (79) TKYGTGKQYLQAIQAASAGMQVYADVFNHKGADSTEWVDAVEVNPNNRQETSCTYQIQAWTKFDFPGRGNTYSSFK
SEQ ID NO.:115 (81) TREGSKHEELVNMTSTAHOYGIKVIADIVINHRAGGDLWNPYVGDYTWTFDFSKVASGKYKAHYMDFHNP-----

161
SEQ ID NO.:113 (156) WHWHYFDGTDWDESRLNRIYKFG--KAWDQEVSNENGNIDYLMYADIDYDHPDVAEEIKRWGTWYANELQLDGFRLLDA
SEQ ID NO.:127 (159) WRWHYFDGTDWDESRLNRIYKFRGTGKAWDQEVDTENGNYDYLMFADLMDHPEVVTTELKNWGTWYVNTTNVDGFRLLDA
SEQ ID NO.:115 (150) ----NYSTSEGTGFGGFPDIDHLPVFNQYWLWASNES-----YAAVLRKSGIDAWRFDY

241
SEQ ID NO.:113 (234) VKHIKESFLRDWVNVHREKTGKEMFTVAEYWQNDLGALENYLNKTNFNHSEVDFVPLHYQFHAASTQGGGYDMRKLNG--
SEQ ID NO.:127 (239) VKHIKYSFFPDWLTHVRSQTRKNLFAYGGEFFSYDVNKLHNYITKTSCTMSLEFQAPLHNNFYTASKSSGYEDMRYLLNN--
SEQ ID NO.:115 (200) VKGYGAWVVKDWLSQWGG-----WAVGEYWDTNVDALLNWAYS--AKVDFPLLYKMDFAFDNKNIPALVYATQNGE

321
SEQ ID NO.:113 (312) TVVSKHPLKAVTFVDNHDTPQGSLSTVQTFKPLAYAFILITRESGYEVEYGYDMYGTGDSQ--REIPALKHKIEPIL
SEQ ID NO.:127 (317) TLMKDQPSLAVTLVDNHDTPQGSLQSWVEPWFKPLAYAFILITRQEGYECVEYGYDYGIPKYN----IPGLKSKIDPLL
SEQ ID NO.:115 (272) TVVSRDPEFKAVTFVANHDTN-----IIWNKYPAFAELITYE-GQEVIFRYDYEEWLNKD-----KLNNL---I

401
SEQ ID NO.:113 (390) KARKQYAYGAQHDYFDHHDIVGWTRREGDSSVANSGLAALITDGPGGAKRMVYGRONAGETWHDITGNRS--EPVVINSEG
SEQ ID NO.:127 (392) IARRDYAYGTORDYIDHODIIGWTRREGIDSKPNSGLAALITDGPGGSKWMYVGGKKGAGKVEYDITGNRS--DVTINADG
SEQ ID NO.:115 (331) WIHEHLAGGSTKILYDDDELIIFMREGYGDRLPL-ITYINLGSDWAEERWVNVGSKFAGYTTHEYTGNLGGWVDRYVQYDG

481
SEQ ID NO.:113 (468) WGEFHVN-----GGSVSIYVQR-----
SEQ ID NO.:127 (470) WGEFKVN-----GGSVSIWVAKTSQVTFVNNATTISGQNVYVVGNIPELGNWNTANAIKMTPSSSYPTWKATIALP
SEQ ID NO.:115 (410) WVKLTAPPHDPANGYYGYSVWSYAGVG-----

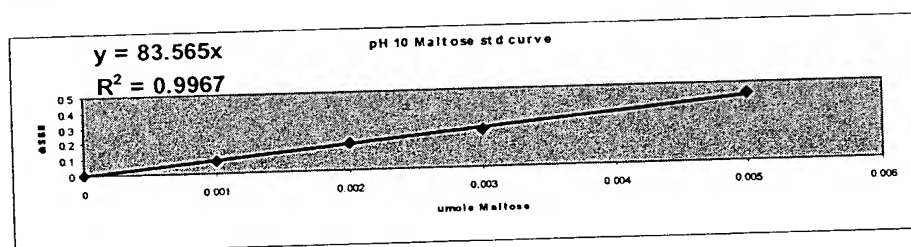
561
SEQ ID NO.:113 (485) -----
SEQ ID NO.:127 (541) QGKAIEFKFIKKDQSGNVVWESIPNRTYTVPLSTGYSYASWNV
SEQ ID NO.:115 (437) -----

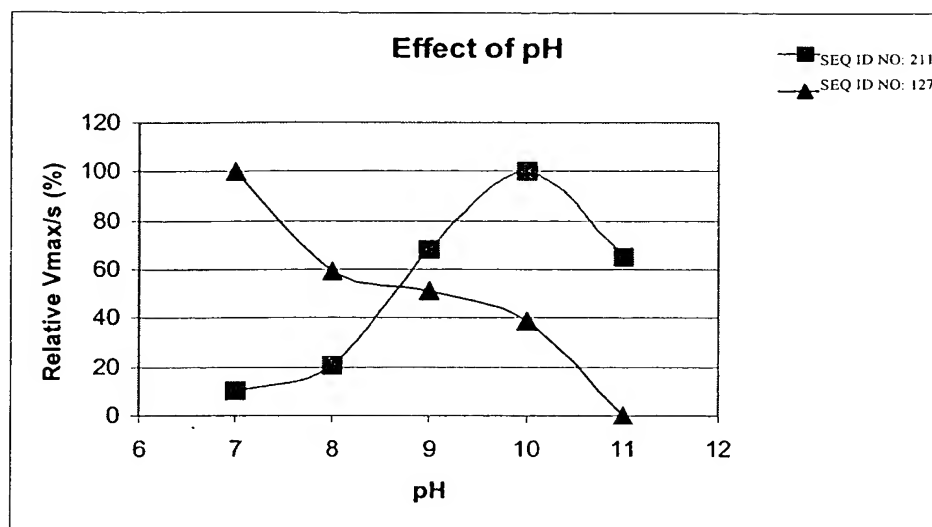
605

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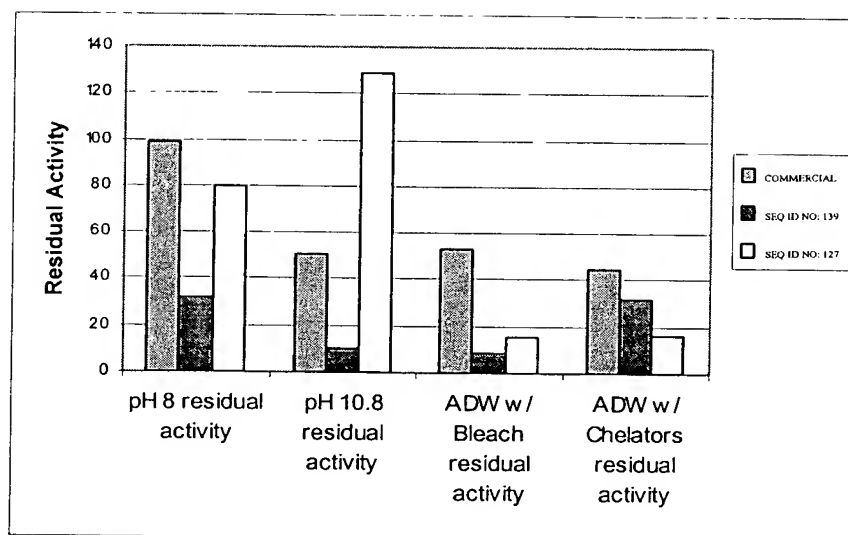
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Figure 11: Example Standard Curve of the assay of Example 5.





**Figure 12:** A graph of the pH rate profiles for 2 different amylases. BD7188 is a control; an enzyme that was discovered previously and has a neutral pH optimum. BD7837 is a more recently discovered amylase and has an optimum around pH 10. Pure protein was used in these assays.



**Figure 13:** Stability of Diversa amylases vs. a commercial enzyme

Figure 14a:

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1
SEQ ID NO: 81 ----- ~~~~~ -MKK FVALFITMFF VVSMAYV... ..AQPASAAK
      pyro ----- ~~~~~ -MKK FVALLITMFF VVSMAAV... ..AQPASAAK
      pyro2 ----- ~~~~~ -VNIKK LTPLLTL LLF FI...VL... ..ASPVSAAK
      thermo SESQCTATCT WRVVYMSAKK LLALLFVLAV LVGVAVIPAR VGIAPVSAGA
      thermo2 ----- ~~~~~ -MA RKVLVALLVF LVVLSVSAVP
      Consensus ----- ~~~~~ -SA-

51
SEQ ID NO: 81 YS..ELEEGG VIMQAFYWDV PGGGIWWDTI RSKIPEWYEA GISAIWIPPA
      pyro YS..ELEEGG VIMQAFYWDV PAGGIWWDTI RSKIPEWYEA GISAIWIPPA
      pyro2 YL..ELEEGG VIMQAFYWDV PGGGIWWDHI RSKIPEWYEA GISAIWLPPP
      thermo TSRPSLEEGG VIMQAFYWDV PAGGIWWDTI RSKIPDWA SA GISAIWIPPA
      thermo2 AKAETLEGG VIMQAFYWDV PGGGIWWDTI AQKIPDWA SA GISAIWIPPA
      Consensus ----- LE-GG VIMQAFYWDV P-GGIWWD-I --KIP-W- -A GISAIW-PP-
              Sense primer

101
SEQ ID NO: 81 SKGMSGGYSM GYDPYDFDL GEYNQKG TIE TRFGSKQELI NMINTAHAYG
      pyro SKGMSGGYSM GYDPYDFDL GEYNQKG TVE TRFGSKQELI NMINTAHAYG
      pyro2 SKGMSGGYSM GYDPYDFDL GEYYQKG TVE TRFGSKKEELV RLIQTAHAYG
      thermo SKGMSGGYSM GYDPYDFDL GEYYQKG TVE TRFGSKQELI NMINTAHAYG
      thermo2 SKGMSGGYSM GYDPYDFDL GEYYQKG SVE TRFGSKKEELV NMINTAHAYN
      Consensus SKGM- G-YSM GYDPYD-FDL GEY-QKG --E TRFGSK-EL- --I-TAH--

151
SEQ ID NO: 81 IKVIADIVIN HRAGGDLEWN PFVGDTY WTD FSKVASGK YT ANYLDFHPNE
      pyro IKVIADIVIN HRAGGDLEWN PFVGDTY WTD FSKVASGK YT ANYLDFHPNE
      pyro2 IKVIADIVIN HRAGGDLEWN PFVGDTY WTD FSKVASGK YT ANYLDFHPNE
      thermo IKVIADIVIN HRAGGDLEWN PFTNSYT WTD FSKVASGK YT ANYLDFHPNE
      thermo2 MKVIADIVIN HRAGGDLEWN PFTNSYT WTD FSKVASGK YT ANYLDFHPNE
      Consensus -KVIAD-VIN HRAGGDLEWN PF---YT WTD FSKVASGK YT ANYLDFHPNE

201
SEQ ID NO: 81 VKCCDEGTFG GFPDIAHEKS WDQHWLW ASD ESYAAYLR SI GVDARWFDY V
      pyro VKCCDEGTFG GFPDIAHEKE WDQHWLW ASD ESYAAYLR SI GVDARWFDY V
      pyro2 LHCCDEGTFG GFPDICHHEK WDQYWLW KSN ESYAAYLR SI GFDGWRFDY V
      thermo VKCCDEGTFG GFPDIAHEKS WDQYWLW ASQ KSYAAYLR SI GIDARWFDY V
      thermo2 LHAGD SGTFG GYPDICHDKS WDQHWLW ASN ESYAAYLR SI GIDARWFDY V
      Consensus ----D-GTFG G-PDI-H-K- WDQ-WLW-S- -SYAAYLR SI G-D-WRFDY V

251
SEQ ID NO: 81 KGYGAWVVKD WLNWWG GWAV GEYWDTN VDA LLNWAYSS GA KVDFDFPLYK
      pyro KGYGAWVVKD WLNWWG GWAV GEYWDTN VDA LLNWAYSS GA KVDFDFPLYK
      pyro2 KGYGAWVVRD WLNWWG GWAV GEYWDTN VDA LLSWAYES GA KVDFDFPLYK
      thermo KGYGAWVVKD WLKWW. ALAV GEYWDTN VDA LLNWAYSS GA KVDFDFPLYK
      thermo2 KGYAP WVVKD WLNWRG GWAV GEYWDTN VDA LLSWAYDS GA KVDFDFPLYK
      Consensus KGY--WVV-- WL--W-- -AV GEYWDTN VDA LL-WAY-S GA KVDFDFPLYK

301
SEQ ID NO: 81 MDEAFDNKNI PALVSA LQNG QTVVSRD PFK AVTFVANH DT DIIWNKYLA Y
      pyro MDEAFDNTNI PALVDA LQNG GTVVSRD PFK AVTFVANH DT DIIWNKYPAY
      pyro2 MDEAFDNNNI PALVYA LQNG QTVVSRD PFK AVTFVANH DT DIIWNKYPAY
      thermo MDEAFDNKNI PALVSA LQNG QTVVSRD PFK AVTFVANH DT DIIWNKYPAY
      thermo2 MDEAFDNNNI PALVDA LKNG GTVVSRD PFK AVTFVANH DT NIIWNKYPAY
      Consensus MDEAFDN-NI PALV-A L-NG -TVVSRD PFK AVTFVANH DT -IIWNKY-AY

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[illegible]

	401				450
SEQ ID NO: 81	MIFVRNGYGS	KPGLITYINL	GSSKVGRWVY	VPKFAGACIH	EYTGNLGGWV
pyro	LIFVRNGDSK	RPGLITYINL	GSSKVGRWVY	VPKFAGACIH	EYTGNLGGWV
pyro2	LIFVRNGDSR	RPGLITYINL	SPNWVGRWVY	VPKFAGACIH	EYTGNLGGWV
thermo	LIFVRNGYGN	KPGLITYINL	GSSKVGRWVY	VPKFAGSCIH	EYTGNLGGWV
thermo2	LIFVRNGYGD	KPGLITYINL	GSSKAGRWVY	VPKFAGSCIH	EYTGNLGGWI
Consensus	-IFVRNG---	-PGLITYINL	-----GRWVY	VPKFAG-CIH	EYTGNLGGW

	451		486
SEQ ID NO: 81	DKYVY SSGWV YFEAPA YDPA	NGQYGY S VWS	YCGVG*
pyro	DKYVE SSGWV YLEAPA YDPA	SGQYGYT VWS	YCGVG*
pyro2	DKRVD SSGWV YLEAPP HDPA	NGYYGY S VWS	YCGVG*
thermo	DKYVG SNGWV YLEAPA HDPA	KGQYGY S VWS	YCGVG*
thermo2	DKWVD SSGRV YLEAPA HDPA	NGQYGY S VWS	YCGVG*
Consensus	DK-V- S-G-V Y-EAP- -DPA	-G-YGY- VWS	YCGVG*
			<b>Antisense primer</b>

Figure 14b:

	1				50
SEQ ID NO: 81	-----	-----	----MKK FVA	LFITMFFV VS	MAVVAQPASA
pyro	-----	-----	----MKK FVA	LLITMFFV VS	MAAVAQPASA
SEQ ID NO: 73	-----	-----	-----	-----	-----
thermo2	-----	-----	-----	-----	-----
SEQ ID NO: 75	-----	-----	-----MA	RKVLVALL VF	LVVLSVSAVP
SEQ ID NO: 77	-----	-----	-----	-----	-----
SEQ ID NO: 83	-----	-----	-----	-----	-----
SEQ ID NO: 85	-----	-----	-----	-----	-----
SEQ ID NO: 79	-----	-----	-----MKP AKL	LVFVLVVS IL	AGLYAQPAGA
thermo	SESQC TATCT	WRVVYM SAKK	LLALLFV LAV	LVGVAVIP AR	VGIAPVSAGA
pyro2	-----	-----	-----VN IKK	LTPLLTL LLF	FIVLASPVSA
CLONE A	-----	-----	-----MRRS ARV	LVLIIAFFLL	AGIYYPSTSA
Consensus	-----	-----	-----	-----	-----
	51				100
SEQ ID NO: 81	AKYSE LEEGG	VIMQAF YWDV	PGGGIWWDTI	RSKIPEWY EA	GISAIWIPPA
pyro	AKYSE LEEGG	VIMQAF YWDV	PAGGIWWDTI	RSKIPEWY EA	GISAIWIPPA
SEQ ID NO: 73	---MA LEEGG	LIMQAF YWDV	PGGGIWWDTI	AQKIPDWA SA	GISAIWIPPA
thermo2	AKAET LENGG	VIMQAF YWDV	PGGGIWWDTI	AQKIPDWA SA	GISAIWIPPA
SEQ ID NO: 75	---MA LEEGG	LIMQAF YWDV	PMGGIWWDTI	AQKIPDWA SA	GISAIWIPPA
SEQ ID NO: 77	---MA LEEGG	LIMQAF YWDV	PMGGIWWDTI	AQKIPDWA SA	GISAIWIPPA
SEQ ID NO: 83	---MA LEEGG	LIMQAF YWDV	PGGGIWWDTI	AQKIPDWA SA	GISAIWIPPA
SEQ ID NO: 85	---MA LEEGG	LIMQAF YWDV	PGGGIWWDTI	AQKIPDWA SA	GISAIWIPPA
SEQ ID NO: 79	AKYLE LEEGG	VIMQAF YWDV	PSGGIWWDTI	RQKIPEWY DA	GISAIWIPPA
thermo	TSRPS LEEGG	VIMQAF YWDV	PAGGIWWDTI	RSKIPDWA SA	GISAIWIPPA
pyro2	AKYLE LEEGG	VIMQAF YWDV	PGGGIWWDTI	RSKIPEWY EA	GISAIWLPPP
CLONE A	AKYSE LEQGG	VIMQAF YWDV	PEGGIWWDTI	RQKIPEWY DA	GISAIWIPPA
Consensus	-----GG	-IMQAF YWDV	P-GGIWWD-I	--KIP-W--A	GISAIW-PP-
	101				150
SEQ ID NO: 81	SKGMS GGYSM	GYDPYD FFDL	GEYNQKG TIE	TRFGSKQE LI	NMINTAHAYG
pyro	SKGMG GAYSM	GYDPYD FFDL	GEYNQKG TVE	TRFGSKQE LI	NMINTAHAYG
SEQ ID NO: 73	SKGMS GGYSM	GYDPYD FFDL	GEYYQKG SVE	TRFGSKKEE LV	NMINTAHAYN
thermo2	SKGMS GGYSM	GYDPYD FFDL	GEYYQKG SVE	TRFGSKKEE LV	NMINTAHAYN
SEQ ID NO: 75	SKGMS GGYSM	GYDPYD YFDL	GEYYQKG TVE	TRFGSKQE LI	NMINTAHAYG
SEQ ID NO: 77	SKGMS GGYSM	GYDPYD YFDL	GEYYQKG TVE	TRFGSKQE LI	NMINTAHAYG
SEQ ID NO: 83	SKGMS GGYSM	GYDPYD FFDL	GEYYQKG TVE	TRFGSKKEE LV	NMINTAHAYG
SEQ ID NO: 85	SKGMS GGYSM	GYDPYD FFDL	GEYYQKG TVE	TRFGSKKEE LV	NMINTAHAYG
SEQ ID NO: 79	SKGMG GAYSM	GYDPYD FFDL	GEYDQKG TVE	TRFGSKQE LV	NMINTAHAYG
thermo	SKGMS GAYSM	GYDPYD FFDL	GEYYQKG TVE	TRFGSKQE LI	NMINTAHAYG
pyro2	SKGMS GGYSM	GYDPYD YFDL	GEYYQKG TVE	TRFGSKKEE LV	RLIQTAHAYG
CLONE A	SKGMG GAYSM	GYDPYD YFDL	GEFYQKG TVE	TRFGSKKEE LV	NMISTAHQYG
Consensus	SKGM- G-YSM	GYDPYD -FDL	GE--QKG --E	TRFGSK-EL-	--I-TAH--
	151				200
SEQ ID NO: 81	IKVIADIVIN	HRAGGD LEWN	PFVGDYT WTD	FSKVASGK YT	ANYLDFHPNE
pyro	IKVIADIVIN	HRAGGD LEWN	PFVGDYT WTD	FSKVASGK YT	ANYLDFHPNE
SEQ ID NO: 73	MKVIADIVIN	HRAGGD LEWN	PFTNSYT WTD	FSKVASGK YT	ANYLDFHPNE
thermo2	MKVIADIVIN	HRAGGD LEWN	PFTNSYT WTD	FSKVASGK YT	ANYLDFHPNE
SEQ ID NO: 75	MKVIADIVIN	HRAGGD LEWN	PFVNDYT WTD	FSKVASGK YT	ANYLDFHPNE
SEQ ID NO: 77	MKVIADIVIN	HRAGGD LEWN	PFVNDYT WTD	FSKVASGK YT	ANYLDFHPNE
SEQ ID NO: 83	IKVIADIVIN	HRAGGD LEWN	PFVNDYT WTD	FSKVASGK YT	ANYLDFHPNE
SEQ ID NO: 85	IKVIADIVIN	HRAGGD LEWN	PFVNDYT WTD	FSKVASGK YT	ANYLDFHPNE
SEQ ID NO: 79	IKVIADIVIN	HRAGGD LEWN	PFVNDYT WTD	FSKVASGK YT	ANYLDFHPNE
thermo	IKVIADIVIN	HRAGGD LEWN	PFTNSYT WTD	FSKVASGK YT	ANYLDFHPNE
pyro2	IKVIADIVIN	HRAGGD LEWN	PFVGDYT WTD	FSKVASGK YT	ANYLDFHPNE
CLONE A	IKVIADIVIN	HRAGGD LEWN	PYVGDYT WTD	FSKVASGK YK	AHYMDFHPPN
Consensus	-KVIAD-VIN	HRAGG- LEWN	P----YT WTD	FSKVASGK Y-	A-Y-DFHPN-

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Figure 14b (cont.)

	201				250
SEQ ID NO: 81	VKCCDEGTFG	GFPDIAHEKS	WDQHWLWASD	ESYAAYLR SI	GVDAWRFDY V
pyro	VKCCDEGTFG	GFPDIAHEKE	WDQHWLWASD	ESYAAYLR SI	GVDAWRFDY V
SEQ ID NO: 73	LHAGD SGTFG	GYPDIC HDKS	WDQHWLWASN	ESYAAYLR SI	GIDAWRFDY V
thermo2	LHAGD SGTFG	GYPDIC HDKS	WDQHWLWASN	ESYAAYLR SI	GIDAWRFDY V
SEQ ID NO: 75	LHAGD SGTFG	GYPDIC HDKS	WDQYWLWASQ	ESYAAYLR SI	GIDAWRFDY V
SEQ ID NO: 77	LHAGD SGTFG	GYPDIC HDKS	WDQYWLWASQ	ESYAAYLR SI	GIDAWRFDY V
SEQ ID NO: 83	LHCCDEGTFG	GYPDIC HDKS	WDQYWLWASS	ESYAAYLR SI	GVDAWRFDY V
SEQ ID NO: 85	LHCCDEGTFG	GYPDIC HDKS	WDQYWLWASS	ESYAAYLR SI	GVDAWCFDY V
SEQ ID NO: 79	VKCCDEGTFG	GFPDIAHEKS	WDQYWLWASN	ESYAAYLR SI	GVDAWRFDY V
thermo	VKCCDEGTFG	GFPDIAHEKS	WDQYWLWASQ	ESYAAYLR SI	GIDAWRFDY V
pyro2	LHCCDEGTFG	GFPDIC HHKE	WDQYWLW KSN	ESYAAYLR SI	GFDGWRFDY V
CLONE A	YSTSD EGTFG	GFPDIDHLVP	FNQYWLWASN	ESYAAYLR SI	GIDAWRFDY V
Consensus	----D-GTFG	G-PDI-H----	--Q-WLW-S-	-SYAAYLR SI	G-D-W-FDY V
	251				300
SEQ ID NO: 81	KGYGA WVVKD	WLNWWG GWAV	GEYWDTN VDA	LLNWAYSS GA	KVFDFPLY Y K
pyro	KGYGA WVVKD	WLNWWG GWAV	GEYWDTN VDA	LLNWAYSS GA	KVFDFPLY Y K
SEQ ID NO: 73	KGYAP WVVKN	WLNWRG GWAV	GEYWDTN VDA	LLSWAYDS GA	KVFDFPLY Y K
thermo2	KGYAP WVVKN	WLNWRG GWAV	GEYWDTN VDA	LLSWAYDS GA	KVFDFPLY Y K
SEQ ID NO: 75	KGYAP WVVVD	WLNWWG GWAV	GEYWDTN VDA	VLNWAYSS GA	KVFDFALY Y K
SEQ ID NO: 77	KGYAP WVVKD	WLNWWG GWAV	GEYWDTN VDA	VLNWAYSS GA	KVFDFALY Y K
SEQ ID NO: 83	KGYGA WVVND	WLSWWG GWAV	GEYWDTN VDA	LLNWAYSS GA	KVFDFPLY Y K
SEQ ID NO: 85	KGYGA WVVND	WLSWWG GWAV	GEYWDTN VDA	LLNWAYNS GA	KVFDFPLY Y K
SEQ ID NO: 79	KGYGA WVVKD	WLDWWG GWAV	GEYWDTN VDA	LLNWAYSS DA	KVFDFPLY Y K
thermo	KGYGA WVVKD	WLKWW. ALAV	GEYWDTN VDA	LLNWAYSS GA	KVFDFPLY Y K
pyro2	KGYGA WVVVD	WLNWWG GWAV	GEYWDTN VDA	LLSWAYES GA	KVFDFPLY Y K
CLONE A	KGYGA WVVKD	WLSQWG GWAV	GEYWDTN VDA	LLNWAYSS GA	KVFDFPLY Y K
Consensus	KGY--WVV--	WL--W---AV	GEYWDTN VDA	-L-WAY-S-A	KVFDF-LYY K
	301				350
SEQ ID NO: 81	MDEAF DNKNI	PALVSA LQNG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYLA Y
pyro	MDEAF DNTNI	PALVDA LQNG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
SEQ ID NO: 73	MDEAF DNNNI	PALVDA LKNG	QTVVSRD PFK	AVTFVANH DT	NIIWNKYPA Y
thermo2	MDEAF DNNNI	PALVDA LKNG	QTVVSRD PFK	AVTFVANH DT	NIIWNKYPA Y
SEQ ID NO: 75	MDEAF DNNNI	PALVDA LRYG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
SEQ ID NO: 77	MDEAF DNNNI	PALVDA LRYG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
SEQ ID NO: 83	MDEAF DNTNI	PALVDA LRYG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
SEQ ID NO: 85	MDEAF DNTNI	PALVYA LKNG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
SEQ ID NO: 79	MDAAF DNKNI	PALVEA LKNG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
thermo	MDEAF DNKNI	PALVSA LQNG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
pyro2	MDEAF DNNNI	PALVYA LQNG	QTVVSRD PFK	AVTFVANH DT	DIIWNKYPA Y
CLONE A	MDEAF DNKNI	PALVYA IQNG	ETVVSRD PFK	AVTFVANH DT	NIIWNKYPA Y
Consensus	MD-AF DN-NI	PALV-A---G	-TVVSRD PFK	AVTFVANH DT	-I IWNKY-AY
	351				400
SEQ ID NO: 81	AFILT YEGQP	VIFYRD YEEW	LNKDRLN NLI	WIHDHLAG GS	TSIVYYDSDE
pyro	AFILT YEGQP	VIFYRD YEEW	LNKDKLN NLI	WIHDHLAG GS	TSIVYYDSDE
SEQ ID NO: 73	AFILT YEGQP	AIFYRD YEEW	LNKDRLR NLI	WIHDHLAG GS	TDI IYYDSDE
thermo2	AFILT YEGQP	AIFYRD YEEW	LNKDRLR NLI	WIHDHLAG GS	TDI IYYDSDE
SEQ ID NO: 75	AFILT YEGQP	TIFYRD YEEW	LNKDKLKNLI	WIHDNLAG GS	TDIVYYDNDE
SEQ ID NO: 77	AFILT YEGQP	TIFYRD YEEW	LNKDKLKNLI	WIHDNLAG GS	TDIVYYDNDE
SEQ ID NO: 83	AFILT YEGQP	VIFYRD YEEW	LNKDKLN NLI	WIHDHLAG GS	TDIVYYDSDE
SEQ ID NO: 85	AFILT YEGQP	VIFYRD YEEW	LNKDKLN NLI	WIHDHLAG GS	TDIVYYDSDE
SEQ ID NO: 79	AFILT YEGQP	TIFYRD YEEW	LNKDRLKNLI	WIHDHLAG GS	TDIVYYDNDE
thermo	AFILT YEGQP	VIFYRD YEEW	LNKDRLKNLI	WIHNNLAG GS	TSIVYYDNDE
pyro2	AFILT YEGQP	VIFYRD FEEW	LNKDKLINLI	WIHDHLAG GS	TTIVYYDNDE
CLONE A	AFILT YEGQP	VIFYRD YEEW	LNKDKLN NLI	WIHEHLAG GS	TKILYYDDDE
Consensus	AFILT YEGQP	-IFYRD -EEW	LNKD-L-NLI	WIH--LAG GS	T-I-YYD-DE

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Figure 14b (cont.)

	401				450
SEQ ID NO: 81	MIFVRNGYGS	KPGLITYINL	GSSKVGRWVY	V.PKFAGACI	HEYTGNLGGW
pyro	LIFVRNGDSK	RPGLITYINL	GSSKVGRWVY	V.PKFAGACI	HEYTGNLGGW
SEQ ID NO: 73	LIFVRNGYGD	KPGLITYINL	GSSKAGR WVY	V.PKFAGSCI	HEYTGNLGGW
thermo2	LIFVRNGYGD	KPGLITYINL	GSSKAGR WVY	V.PKFAGSCI	HEYTGNLGGW
SEQ ID NO: 75	LIFVRNGYGS	KPGLITYINL	GSSKAGR WVY	V.PKFAGSCI	HEYTGNLGGW
SEQ ID NO: 77	LIFVRNGYGS	KPGLITYINL	ASSKAGR WVY	V.PKFAGSCI	HEYTGNLGGW
SEQ ID NO: 83	LIFVRNGYGT	KPGLITYINL	GSSKVGR WVY	V.PKFAGSCI	HEYTGNLGGW
SEQ ID NO: 85	LIFVRNGYGT	KPGLITYINL	GSSKAGR WVY	V.PKFAGSCI	HEYTGNLGGW
SEQ ID NO: 79	LIFVRNGYGD	KPGLITYINL	GSSKAGR WVY	V.PKFAGACI	HEYTGNLGGW
thermo	LIFVRNGYGN	KPGLITYINL	GSSKVGR WVY	V.PKFAGSCI	HEYTGNLGGW
pyro2	LIFVRNGDSR	RPGLITYINL	SPNWVGR WVY	V.PKFAGACI	HEYTGNLGGW
CLONE A	LIFMR EGYGD	RPGLITYINL	GSDWAER WVN	VGSKFAGYTI	HEYTGNLGGW
Consensus	-IF-R-G---	-PGLITYINL	-----RWV-	V--KFAG--I	HEYTG-LGGW

	451				487
SEQ ID NO: 81	VDKYV YSSGW	VYFEAP AYDP	ANGQYGY SVW	SYCGVG*	
pyro	VDKYV ESSGW	VYLEAP AYDP	ASGQYGY TVW	SYCGVG*	
SEQ ID NO: 73	IDKWV DSSGR	VYLEAP AHDP	ANGQYGY SVW	SYCGVG*	
thermo2	IDKWV DSSGR	VYLEAP AHDP	ANGQYGY SVW	SYCGVG*	
SEQ ID NO: 75	VDKWV DSSGW	VYLEAP AHDP	ANGQYGY SVW	SYCGVG*	
SEQ ID NO: 77	VDKWV DSSGW	VYLEAP AHDP	ANGQYGY SVW	SYCGVG*	
SEQ ID NO: 83	IDKYV SSSGW	VYLEAP AHDP	ANGYYGY SVW	SYCGVG*	
SEQ ID NO: 85	IDKYV SSSGW	VYLEAP AHDP	ANGQYGY SVW	SYCGVG*	
SEQ ID NO: 79	VDKWV DSSGW	VYLEAP AHDP	ANGYYGY SVW	SYCGVG*	
thermo	VDKYV GSNW	VYLEAP AHDP	AKGQYGY SVW	SYCGVG*	
pyro2	VDKRV DSSGW	VYLEAP PHDP	ANGYYGY SVW	SYCGVG*	
CLONE A	VDRYV QYDGW	VKLTAP PHDP	ANGYYGY SVW	SYAGVG*	
Consensus	-D--V---G-	V---AP--DP	A-G-YGY -VW	SY-GVG*	

Figure 14c:

	1				50
SEQ ID NO: 83	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
SEQ ID NO: 85	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
SEQ ID NO: 75	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
SEQ ID NO: 77	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
SEQ ID NO: 73	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
SEQ ID NO: 79	~~~ATGA AGC	CTGCGAAA CT	CCTCGTCTT T	GTGCTCGTAG	TCTCTATCCT
SEQ ID NO: 81	~~~ATGA AGA	AGTTTGTC GC	CCTGTTCAT A	ACCATGTTTT	TCGTAGTGAG
CLONE A	ATGAGGA GAT	CCGCAAGG GT	TTTGTTCT G	ATTATAGCGT	TTTTCTCTCT
Consensus	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
	51				100
SEQ ID NO: 83	~~~~~	~~~~~	~~~~~	~~~~~	ATGGCTCTGG
SEQ ID NO: 85	~~~~~	~~~~~	~~~~~	~~~~~	ATGGCTCTGG
SEQ ID NO: 75	~~~~~	~~~~~	~~~~~	~~~~~	ATGGCTCTGG
SEQ ID NO: 77	~~~~~	~~~~~	~~~~~	~~~~~	ATGGCTCTGG
SEQ ID NO: 73	~~~~~	~~~~~	~~~~~	~~~~~	ATGGCTCTGG
SEQ ID NO: 79	CGCGGGG CTC	TACGCCCCA GC	CCGCGGGGG C	GGCCAAGTAC	CTGGAGCTCG
SEQ ID NO: 81	CATGGCA GTC	GTTGCACAG C	CAGCTAGCG C	CGCAAAGTAT	TCCGAGCTCG
CLONE A	GGCGGGG ATT	TACTACCC CT	CCACGAGTG C	CGCGAAGTAC	TCCGAGCTGG
Consensus	~~~~~	~~~~~	~~~~~	~~~~~	~~~~~
	101				150
SEQ ID NO: 83	AAGAGGG CGG	GCTCATAA TG	CAGGCCTTC T	ACTGGGATGT	TCCTGGAGGA
SEQ ID NO: 85	AAGAGGG CGG	GCTTATAA TG	CAGGCATTC T	ATTGGGACGT	CCCAGGTGGA
SEQ ID NO: 75	AAGAGGG CGG	GCTTATAA TG	CAGGCATTC T	ACTGGGACGT	CCCCATGGGA
SEQ ID NO: 77	AAGAGGG CGG	GCTCATAA TG	CAGGCCTTC T	ACTGGGACGT	CCCCATGGGA
SEQ ID NO: 73	TAGAGGG CGG	GCTTATAA TG	CAGGCCTTC T	ACTGGGACGT	CCCAGGTGGA
SEQ ID NO: 79	AAGAGGG CGG	CGTCATAA TG	CAGGCGTTC T	ACTGGGACGT	GCCTTCAGGA
SEQ ID NO: 81	AAGAAGG CGG	CGTTATAA TG	CAGGCCTTC T	ACTGGGACGT	CCCAGGTGGA
CLONE A	AGCAGGG CGG	AGTCATAA TG	CAGGCCTTC T	ACTGGGACGT	TCCGAGGGGA
Consensus	-----GG CGG	--T-ATAA TG	CAGGC-TTC T	A-TGGGA-GT	-CC----GGA
	151				200
SEQ ID NO: 83	GGAATCT GGT	GGGACACA AT	AGCTCAAAA G	ATACCCGAAT	GGGCAAGTGC
SEQ ID NO: 85	GGAATCT GGT	GGGACACC AT	AGCCCAGAA G	ATACCCGAAT	GGGCAAGTGC
SEQ ID NO: 75	GGAATCT GGT	GGGACACG AT	AGCCCAGAA G	ATACCCGACT	GGGCAAGCGC
SEQ ID NO: 77	GGAATCT GGT	GGGACACG AT	AGCCCAGAA G	ATACCCGACT	GGGCAAGCGC
SEQ ID NO: 73	GGAATCT GGT	GGGACACC AT	AGCCCAGAA G	ATACCCGACT	GGGCGAGCGC
SEQ ID NO: 79	GGAATAT GGT	GGGACACA AT	ACGGCAGAA G	ATACCGGAGT	GGTACGATGC
SEQ ID NO: 81	GGAATCT GGT	GGGACACC AT	CAGGAGCAA G	ATACCGGAGT	GGTACGAGGC
CLONE A	GGAATCT GGT	GGGACACA AT	ACGGCAGAA G	ATCCCTGAAT	GGTACGATGC
Consensus	GGAAT-T GGT	GGGACAC- AT	-----AA G	AT-CC-GA-T	GG-----GC
	201				250
SEQ ID NO: 83	AGGAATC TCA	GCGATATG GA	TTCCACCAG C	GAGTAAGGGC	ATGAGCGGTG
SEQ ID NO: 85	AGGAATC TCA	GCGATATG GA	TTCCACCAG C	GAGTAAGGGA	ATGAGCGGTG
SEQ ID NO: 75	CGGGATT TCG	GCGATATG GA	TTCCCCCG C	GAGCAAGGGT	ATGAGCGGCG
SEQ ID NO: 77	CGGGATT TCG	GCGATATG GA	TCCCTCCCG C	GAGCAAGGGT	ATGAGCGGCG
SEQ ID NO: 73	CGGGATT TCG	GCAATATG GA	TTCTCCCG C	GAGTAAGGGC	ATGAGCGGCG
SEQ ID NO: 79	CGGAATC TCC	GCAATATG GA	TTCCCCCG C	GAGCAAGGGC	ATGGGCGGCG
SEQ ID NO: 81	GGGAATA TCC	GCCATTG GA	TTCCGCCAG C	CAGCAAGGGG	ATGAGCGGCG
CLONE A	AGGCATA TCC	GCCATCTG GA	TACCCCGG C	GAGCAAGGGC	ATGGGCGGGG
Consensus	-GG-AT- TC-	GC-AT-TG GA	T-CC-CC-G C	-AG-AAGGG-	ATG-GCGG-G

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Figure 14c (cont.)

	251		300
SEQ ID NO: 83	GTTATTC CAT	GGGCTACG AT	CCCTACGAT T
SEQ ID NO: 85	GTTATTC CAT	GGGCTACG AT	CCCTACGAT T
SEQ ID NO: 75	GCTATTC GAT	GGGCTACG AC	CCCTACGAT T
SEQ ID NO: 77	GCTATTC GAT	GGGCTACG AC	CCCTACGAT T
SEQ ID NO: 73	GCTATTC GAT	GGGCTACG AC	CCCTACGAT T
SEQ ID NO: 79	CCTATTC GAT	GGGCTACG AC	CCCTACGAT T
SEQ ID NO: 81	GTTACTC GAT	GGGCTACG AT	CCCTACGAT T
CLONE A	CCTACTC GAT	GGGCTACG AC	CCCTACGAT T
Consensus	--TA-TC-AT	GGGCTACG A-	CCCTACGA- T

	301		350
SEQ ID NO: 83	TATCAGA AGG	GGACAGTT GA	GACGCGCTT C
SEQ ID NO: 85	TATCAGA AGG	GGACAGTT GA	GACGCGCTT C
SEQ ID NO: 75	TACCAGA AGG	GAACGGTG GA	AACAAGATT C
SEQ ID NO: 77	TACCAGA AGG	GAACGGTG GA	AACGAGGTT C
SEQ ID NO: 73	TACCAGA AGG	GAACGGTG GA	AACGAGGTT C
SEQ ID NO: 79	GACCAGA AGG	GAACGGTA GA	GACCGCTT C
SEQ ID NO: 81	AACCAGA AGG	GAACCATC GA	AACGCGCTT T
CLONE A	TACCAGA AGG	GAACGGTG GA	GACCGCTT C
Consensus	-A-CAGA AGG	G-A--T- GA	-AC--G-TT -

	351		400
SEQ ID NO: 83	GAACATG ATA	AACACCGC AC	ACTCCTACG G
SEQ ID NO: 85	GAACATG ATA	AACACCGC AC	ACTCCTACG G
SEQ ID NO: 75	AAACATG ATA	AACACCGC CC	ACGCTATG G
SEQ ID NO: 77	AAACATG ATA	AACACCGC CC	ACGCTATG G
SEQ ID NO: 73	GAACATG ATA	AACACCGC CC	ATGCTCACA A
SEQ ID NO: 79	GAACATG ATA	AACACCGC CC	ACGCTACG G
SEQ ID NO: 81	CAATATG ATA	AACACGGC CC	ATGCCTACG G
CLONE A	CAACATG ATC	TCCACGGC CC	ACCAATACG G
Consensus	-AA-ATG AT-	--CAC-GC -C	A-----A-- -

	401		450
SEQ ID NO: 83	TAGTCAT AAA	CCACCGCG CC	GGTGGAGAC C
SEQ ID NO: 85	TAGTCAT AAA	CCACCGCG CC	GGTGGAGGC C
SEQ ID NO: 75	TAGTCAT CAA	CCACCGCG CC	GGCGGCGAT C
SEQ ID NO: 77	TAGTCAT CAA	CCACCGCG CC	GGCGGCGAC C
SEQ ID NO: 73	TAGTCAT CAA	CCACCGCG CC	GGCGGCGAC C
SEQ ID NO: 79	TAGTAAT CAA	CCACCGCG CC	GGAGGAGAC C
SEQ ID NO: 81	TCGTCAT AAA	CCACCGCG CA	GGCGGAGAC C
CLONE A	TAGTGAT AAA	CCACCGCG CA	GGTGGAGAC C
Consensus	T-GT-AT -AA	CCACCGCG C-	GG-GG-G-- C

	451		500
SEQ ID NO: 83	AACGACT ATA	CCTGGACA GA	CTTCTCAAAA A
SEQ ID NO: 85	AACGACT ATA	CCTGGACA GA	CTTCTCAAAA A
SEQ ID NO: 75	AACGACT ATA	CCTGGACC GA	CTTCTCGAA G
SEQ ID NO: 77	AACGACT ATA	CCTGGACC GA	CTTCTCAAA G
SEQ ID NO: 73	AACAGCT ACA	CCTGGACC GA	TTTCTCGAA G
SEQ ID NO: 79	AATGACT ACA	CCTGGACG GA	CTTCTCGAA G
SEQ ID NO: 81	GGGGACT ACA	CCTGGACG GA	CTTCTCAAA G
CLONE A	GGCGACT ATA	CCTGGACG GA	CTTTTCTAA G
Consensus	-----CT A-A	CCTGGAC- GA	-TT-TC-AA -

Figure 14c (cont.)

	501		550
SEQ ID NO: 83	GGCCAAC TAC CTTGACTT CC ACCCAAACG A GCTTCACTGT TGTGATGAAG		
SEQ ID NO: 85	AGCCAAC TAC CTTGACTT CC ACCCAAACG A GCTTCACTGT TGTGATGAAG		
SEQ ID NO: 75	GGCCAAC TAC CTCGACTT CC ACCCGAACG A GCTCCACGCG GGCGATTCCG		
SEQ ID NO: 77	GGCCAAC TAC CTCGACTT CC ACCCGAACG A GCTCCATGCG GGCGATTCCG		
SEQ ID NO: 73	GGCCAAC TAC CTCGACTT CC ACCCGAACG A GCTTCACTGT GGCGATTCCG		
SEQ ID NO: 79	GGCCAAC TAC CTCGACTT CC ACCCAAACG A GGTCAAGTGC TGGGACGAGG		
SEQ ID NO: 81	TGCCAAC TAC CTCGACTT CC ACCCAAACG A GGTCAAGTGC TGTGACGAGG		
CLONE A	GGCCAC TAC ATGGACTT CC ATCCAAACA A CTACAGCACC TCAGACGAGG		
Consensus	-GCC-AC TAC -T-GACTT CC A-CC-AAC- A ----- ---GA----		
	551		600
SEQ ID NO: 83	GTACCTT TGG AGGATACC CT GATATATGT C ACGACAAAAG CTGGGACCAG		
SEQ ID NO: 85	GTACCTT TGG AGGATACC CT GATATATGT C ACGACAAAAG CTGGGACCAG		
SEQ ID NO: 75	GAACATT TGG AGGCTATC CC GACATATGC C ACGACAAGAG CTGGGACCAG		
SEQ ID NO: 77	GAACATT TGG AGGCTATC CC GACATATGC C ACGACAAGAG CTGGGACCAG		
SEQ ID NO: 73	GAACATT TGG AGGCTATC CC GACATATGC C ACGACAAGAG CTGGGACCAG		
SEQ ID NO: 79	GCACCTT TGG AGGGTTC C GACATAGCC C ACGAGAAGAG CTGGGACCAG		
SEQ ID NO: 81	GCACATT TGG AGGCTTC CA GACATAGCC C ACGAGAAGAG CTGGGACCAG		
CLONE A	GAACCTT CGG TGGCTTC CA GACATTGAT C ACCTCGTGCC CTTCAACCAG		
Consensus	G-AC-TT -GG -GG-T--C C- GA-AT---- C AC----- CT---ACCAG		
	601		650
SEQ ID NO: 83	TACTGGC TCT GGGCGAGC AG CGAAAGCTA C GCTGCCTACC TCAGGAGCAT		
SEQ ID NO: 85	TACTGGC TCT GGGCGAGC AG CGAAAGCTA C GCTGCCTACC TCAGGAGCAT		
SEQ ID NO: 75	TACTGGC TCT GGGCCAGC CA GGAGAGCTA C GCGGCCTATC TCAGGAGCAT		
SEQ ID NO: 77	TACTGGC TCT GGGCCAGC CA GGAGAGCTA C GCGGCATATC TCAGGAGCAT		
SEQ ID NO: 73	CACTGGC TCT GGGCCAGC AA CGAAAGCTA C GCCGCCTACC TCCGGAGCAT		
SEQ ID NO: 79	TACTGGC TCT GGGCGAGC AA CGAGAGCTA C GCCGCCTACC TCAGGAGCAT		
SEQ ID NO: 81	CACTGGC TCT GGGCGAGC GA TGAGAGCTA C GCCGCCTACC TAAGGAGCAT		
CLONE A	TACTGGC TGT GGGCGAGC AA CGAGAGCTA C GCCGCCTACC TCAGGAGCAT		
Consensus	-ACTGGC T-T GGGC-AGC -- -GA-AGCTA C GC-GC-TA-C T--GGAGCAT		
	651		700
SEQ ID NO: 83	AGGGGTT GAC GCCTGGCG TT TCGACTACG T CAAGGGCTAC GGAGCATGGG		
SEQ ID NO: 85	AGGGGTT GAC GCCTGGTG TT TCGACTACG T CAAGGGCTAC GGAGCCTGGG		
SEQ ID NO: 75	CGGCATC GAC GCCTGGCG CT TCGACTACG T CAAGGGCTAT GCTCCCTGGG		
SEQ ID NO: 77	CGGCATC GAT GCCTGGCG CT TCGACTACG T CAAGGGCTAT GCTCCCTGGG		
SEQ ID NO: 73	CGGCATC GAC GCCTGGCG CT TCGACTACG T CAAGGGCTAC GCTCCCTGGG		
SEQ ID NO: 79	CGGCGTT GAC GCATGGCG CT TCGACTACG T CAAGGGCTAC GGAGCGTGGG		
SEQ ID NO: 81	CGGCGTT GAT GCCTGGCG CT TTGACTACG T GAAGGGCTAC GGAGCGTGGG		
CLONE A	AGGGATC GAT GCGTGGCG CT TTGACTACG T TAAGGGCTAC GCGCGTGGG		
Consensus	-GG--T- GA- GC-TGG-G -T T-GACTACG T -AAGGGCTA- G---C-TGGG		
	701		750
SEQ ID NO: 83	TTGTAA CGA CTGGCTCA GC TGGTGGGGA G GCTGGGCCGT TGGAGAGTAC		
SEQ ID NO: 85	TTGTAA CGA CTGGCTCA GC TGGTGGGGA G GCTGGGCCGT TGGAGAGTAC		
SEQ ID NO: 75	TCGTCA GGA CTGGCTGA AC TGGTGGGGA G GCTGGGCGT TGGAGAGTAC		
SEQ ID NO: 77	TCGTCA GGA CTGGCTGA AC TGGTGGGGA G GCTGGGCGT TGGAGAGTAC		
SEQ ID NO: 73	TCGTAA GAA CTGGCTGA AC CGGTGGGGA G GCTGGGCGT TGGAGAGTAC		
SEQ ID NO: 79	TCGTCA GGA CTGGCTGA AC TGGTGGGGA G GCTGGGCCGT CGGGAGTAC		
SEQ ID NO: 81	TCGTCA GGA CTGGCTCA AC TGGTGGGGA G GCTGGGCCGT TGGAGTAC		
CLONE A	TCGTCA GGA CTGGCTGA GT CAGTGGGGA G GCTGGGCCGT CGGCGAGTAC		
Consensus	T-GT-A- --A CTGGCT- - - -GTGGGGA G GCTGGGC-GT -GG-GAGTAC		

Figure 14c (cont.)

	751				800
SEQ ID NO: 83	TGGGACA CGA	ACGTTGAT GC	ACTCCTCAA C	TGGGCATACA	GCAGCGGCGC
SEQ ID NO: 85	TGGGACA CTA	ACGTTGAT GC	ACTCCTCAA C	TGGGCATACA	ACAGCGGCGC
SEQ ID NO: 75	TGGGACA CCA	ACGTTCGAC GC	TGTTCTCAA C	TGGGCATACT	CGAGCGGTGC
SEQ ID NO: 77	TGGGACA CCA	ACGTTCGAC GC	TGTTCTCAA C	TGGGCATACT	CGAGCGGTGC
SEQ ID NO: 73	TGGGACA CCA	ACGTTCGAT GC	ACTCCTGAG C	TGGGCCTACG	ACAGCGGTGC
SEQ ID NO: 79	TGGGACA CAA	ACGTTGAT GC	ACTGCTCAA C	TGGGCCTACT	CGAGCGATGC
SEQ ID NO: 81	TGGGACA CCA	ACGTTGAT GC	ACTCCTCAA C	TGGGCCTACT	CGAGCGGCGC
CLONE A	TGGGACA CCA	ACGTTCGAT GC	GCTCCTCAA C	TGGGCCTACA	GCAGCGGCGC
Consensus	TGGGACA C-A	ACGT-GA-GC	--T-CT-A-C	TGGGC-TAC-	--AGCG--GC
	801				850
SEQ ID NO: 83	CAAGGTC TTT	GACTTCCC GC	TCTACTACA A	GATGGACGAA	GCCTTCGACA
SEQ ID NO: 85	CAAGGTC TTT	GACTTCCC GC	TCTACTACA A	GATGGACGAA	GCCTTCGACA
SEQ ID NO: 75	CAAGGTC TTT	GACTTCGC CC	TCTACTACA A	GATGGACGAG	GCCTTCGATA
SEQ ID NO: 77	CAAGGTC TTT	GACTTCGC CC	TCTACTACA A	GATGGACGAG	GCCTTCGATA
SEQ ID NO: 73	TAAAGTC TTC	GACTTCCC GC	TCTACTACA A	GATGGACGAG	GCCTTCGATA
SEQ ID NO: 79	AAAAGTC TTC	GACTTCCC GC	TCTACTACA A	GATGGACGCG	GCCTTTGACA
SEQ ID NO: 81	CAAGGTC TTC	GACTTCCC GC	TCTACTACA A	GATGGATGAG	GCCTTTGACA
CLONE A	CAAGGTC TTC	GACTTCCC GC	TCTACTACA A	GATGGACGAG	GCCTTTGACA
Consensus	-AA-GTC TT-	GACTTC-C-C	TCTACTACA A	GATGGA-G--	GCCTT-GA-A
	851				900
SEQ ID NO: 83	ACACCAA CAT	CCCGGCAT TA	GTGGATGCA C	TCAGATACGG	CCAGACAGTG
SEQ ID NO: 85	ATACCAA CAT	CCCGCTT TG	GTTTACGCC C	TCAAGAATGG	CGGGACAGTG
SEQ ID NO: 75	ACAACAA CAT	TCCCGCCC TG	GTGGACGCC C	TCAGATACGG	CCAGACAGTG
SEQ ID NO: 77	ACAACAA CAT	TCCCGCCC TG	GTGGACGCC C	TCAGATACGG	TCAGACAGTG
SEQ ID NO: 73	ACAACAA CAT	CCCGGCC TC	GTGGACGCC C	TCAAGAACGG	AGGCACGGTC
SEQ ID NO: 79	ACAAGAA CAT	TCCCGCAC TC	GTGGAGGCC C	TCAAGAACGG	GGGCACAGTC
SEQ ID NO: 81	ACAAAAA CAT	TCCAGCGC TC	GTCTCTGCC C	TTCAGAACGG	CCAGACTGTT
CLONE A	ACAAGAA CAT	TCCCGCCC TC	GTTTACGCC A	TCCAGAACGG	TGAAACCGTC
Consensus	A-A--AA CAT	-CC-GC--T-	GT----GC--	T-----A-GG	----AC-GT-
	901				950
SEQ ID NO: 83	GTCAGCC GCG	ATCCCTTC AA	GGCGGTAAC T	TTCGTTGCCA	ACCACGATAC
SEQ ID NO: 85	GTCAGCC GCG	ACCCATTC AA	GGCGGTAAC T	TTCGTTGCCA	ACCACGATAC
SEQ ID NO: 75	GTCAGCC GCG	ACCCGTTT AA	GGCTGTGAC G	TTGTAGCCA	ACCACGATAC
SEQ ID NO: 77	GTCAGCC GCG	ACCCGTTT AA	GGCTGTGAC G	TTGTAGCCA	ACCACGATAC
SEQ ID NO: 73	GTCAGCC GCG	ACCCGTTT AA	AGCCGTGAC C	TTCGTTGCCA	ACCACGATAC
SEQ ID NO: 79	GTCAGCC GCG	ACCCGTTT AA	GGCCGTAAC C	TTCGTTGCCA	ACCACGACAC
SEQ ID NO: 81	GTCTCCC GCG	ACCCGTTT AA	GGCCGTAAC C	TTGTAGCAA	ACCACGACAC
CLONE A	GTCAGCA GGG	ATCCCTTC AA	GGCCGTTAC C	TTCGTTGGCTA	ACCACGATAC
Consensus	GTC--C-G-G	A-CC-TT-AA	-GC-GT-AC-	TT-GT-GC-A	ACCACGA-AC
	951				1000
SEQ ID NO: 83	AGATATA ATC	TGGAACAA GT	ATCCGGCTT A	TGCATTTCATC	CTTACCTATG
SEQ ID NO: 85	AGATATA ATC	TGGAACAA GT	ATCCGGCTT A	TGCATTTCATC	CTTACCTATG
SEQ ID NO: 75	CGACATA ATC	TGGAACAA GT	ATCCAGCCT A	CGCGTTTCATC	CTCACCTACG
SEQ ID NO: 77	CGACATA ATC	TGGAACAA GT	ATCCAGCCT A	CGCGTTTCATC	CTCACCTACG
SEQ ID NO: 73	CAACATA ATC	TGGAACAA GT	ATCCGGCCT A	CGCCTTCATC	CTCACCTATG
SEQ ID NO: 79	GGACATA ATT	TGGAACAA GT	ACCCGGCCT A	CGCCTTCATC	CTCACCTACG
SEQ ID NO: 81	CGATATA ATC	TGGAACAA GT	ACCTTGCTT A	TGCTTTTCATC	CTCACCTACG
CLONE A	GAACATA ATC	TGGAACAA GT	ACCCTGCCT A	TGCCTTCATC	CTGACCTACG
Consensus	--A-ATA AT-	TGGAACAA GT	A-C--GC-TA	-GC-TTCATC	CT-ACCTA-G



Figure 14c (cont.)

	1251				1300
SEQ ID NO: 83	CCACGAG TAC	ACCGGCAA CC	TCGGCGGTT G	GATAGACAAG	TACGTCTCCT
SEQ ID NO: 85	CCACGAG TAC	ACCGGCAG CC	TCGGCGGTT G	GATAGACAAG	TACGTCTCCT
SEQ ID NO: 75	ACACGAG TAC	ACCGGCAA CC	TCGGCGGCT G	GGTGGACAAG	TGGGTGGACT
SEQ ID NO: 77	ACACGAG TAC	ACCGGCAA TC	TCGGCGGCT G	GGTGGACAAG	TGGGTGGACT
SEQ ID NO: 73	ACACGAG TAC	ACCGGCAA CC	TCGGCGGCT G	GATTGACAAG	TGGGTGGACT
SEQ ID NO: 79	CCACGAG TAC	ACCGGCAA CC	TCGGCGGCT G	GGTGGACAAG	TGGGTGGACT
SEQ ID NO: 81	CCACGAG TAT	ACTGGTAA CC	TCGGAGGCT G	GGTAGACAAG	TACGTCTACT
CLONE A	CCACGAA TAC	ACCGGAAA CC	TCGGCGGCT G	GGTCGACAGG	TACGTCCAGT
Consensus	-CACGA- TA-	AC-GG-A- -C	TCGG-GG-TG	G-T-GACA-G	T--GT----T

	1301				1350
SEQ ID NO: 83	CCAGCGG CTG	GGTCTATC TT	GAGGCCCCAG	CCCACGACCC	GGCGAACGGC
SEQ ID NO: 85	CCAGCGG CTG	GGTCTACC TT	GAGGCCCCG G	CCCACGACCC	GGCCAATGGC
SEQ ID NO: 75	CAAGCGG CTG	GGTTTACC TC	GAGGCTCCT G	CCCACGACCC	GGCCAACGGC
SEQ ID NO: 77	CAAGCGG CTG	GGTCTACC TC	GAGGCTCCT G	CCCACGACCC	GGCCAACGGC
SEQ ID NO: 73	CAAGCGG TCG	GGTCTACC TT	GAGGCCCCG G	CCCACGACCC	GGCCAACGGC
SEQ ID NO: 79	CAAGCGG GTG	GGTGTACC TC	GAGGCCCCCT G	CCCACGACCC	GGCCAACGGC
SEQ ID NO: 81	CAAGCGG CTG	GGTCTATT TC	GAAGCTCCA G	CTTACGACCC	TGCCAACGGG
CLONE A	ACGACGG CTG	GGTCAAGC TT	ACCGCTCCG C	CACACGATCC	GGCAAACGGC
Consensus	----CGG --G	GGT--A-- T-	---GC-CC- -	C--ACGA-CC	-GC-AA-GG-

	1351				1393
SEQ ID NO: 83	TACTACG GCT	ACTCCGTA TG	GAGCTACTG C	GGGGTTGGGT	GA~
SEQ ID NO: 85	CAGTATG GCT	ACTCCGTC TG	GAGCTATTG C	GGGGTTGGGT	GA~
SEQ ID NO: 75	CAGTACG GCT	ACTCCGTT TG	GAGCTATTG C	GGTGTGGGT	GA~
SEQ ID NO: 77	CAGTACG GCT	ACTCCGTC TG	GAGCTACTG C	GGTGTGGGT	GA~
SEQ ID NO: 73	CAGTACG GCT	ACTCCGTA TG	GAGCTACTG C	GGTGTGGGT	GA~
SEQ ID NO: 79	TATTACG GCT	ACTCCGTC TG	GAGCTACTG C	GGGGTGGGCT	GA~
SEQ ID NO: 81	CAGTATG GCT	ACTCCGTG TG	GAGCTATTG C	GGTGTGGGT	GA~
CLONE A	TATTACG GCT	ACTCCGTC TG	GAGCTACGC C	GGAGTTGGAT	GA~
Consensus	-A-TA-G GCT	ACTC-GT- TG	GAGCTA-- C	GG-GT-GG-T	GA~

Figure 15:

# Neighbor-joining tree for Thermococcales

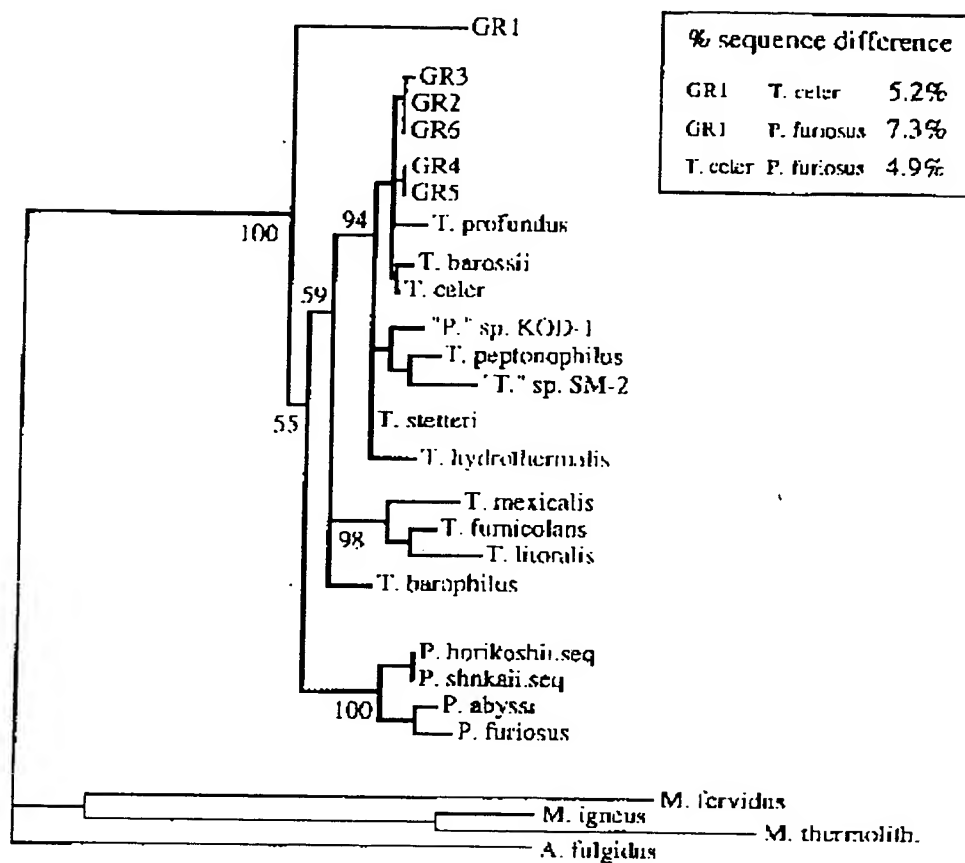


Figure 16 (all sequences are listed in 5' to 3' order)

SEQ ID NO.: 1

atggcaaagtattccgagctcgaagagggcgggctcataatgcaggccttctactgggacgtcccatgggaggaatctggtgggacacgat  
agcccagaagatacccgactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcatgggcggcgctattcgatg  
ggctacgacccctacgacttcttgacctcggtagtacgaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacaccgcccacgcctatggcatgaaggtaatagccgatatagtcatcaaccaccgcggcggtgacctggagtggaaaccctt  
cgtgaacgactatacctggaccgacttctcaaaggctgcgtcgggtaaatacacggccaactacctcgaactccaccgaacgagctccatgc  
ggcgattccggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggcccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgatgcctggcgcttcgactacgtcaagggtctacggagcgtgggtcgtcaaggactggctggactg  
gtggggaggtgggctggcggtcggggagtactgggacacaaacgttgatgactgctcaactgggctactcagcgtatgcaaaagtcttcgactt  
cccgctctactacaagatggacgcggcctttgacaacaagaacattcccgcactcgtcaggccctcaagaacgggggacagctcgtcagcc  
gcgacccgtttaaggcgttaaccttcgttgcaaacacgacaccgatataatctggaacaagtatccagcctacgcgttcacctcacctacgag  
ggccagccgacaatattctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccgg  
aggaagcactgacatcgtttactacgacaacgacgagctgataatcgtgagaaacggctacggaagcaagccgggactgataacatacatcaa  
cctcgctcaagcaagccggaagggtgggtttacgttcggaagtcgcaggctcgtgcatacacgagtacaccggcaatctcggcggctgggt  
ggacaagtgggtgactcaagcggtgggtctacctcaggctcgtgccacgaccggccaacggccagtacgggtactccgtctggagc  
tactgcggtgttgggtga

SEQ ID NO.: 2

Met Ala Lys Tyr Ser Glu Leu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val  
Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Ala  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 3

atggccaagtacctggagctcgaagagggcgggctcataatgcaggccttctactgggacgtcccatgggaggaatctggtgggacacgat  
agcccagaagatacccgactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcatgggcggcgctattcgatg  
ggctacgacccctacgacttcttgacctcggtagtacgaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacaccgcccacgcctacggcatcaaggctatcgcagacatagtaatcaaccaccgcggcgaggagaccttgagtggaaaccct  
tcgtcaatgactacacctggacggacttctcgaaggctcgttcggcgaagtacacggccaattacctcgaactccaccgaacgagctccatgc  
ggcgattccggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggcccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgatgcctggcgcttcgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactggt  
ggggaggtcggcggttggagagtactgggacaccaacgtcgacgtgttctcaactgggcatactcagcgggtgccaaggtctttgacttcg  
ccctctactacaagatggatgaggcctttgacaacaaaaacattccagcgtcgtctcgtcccttcagaacggccagactgttctcccgac  
ccgttcaaggccgtaacctttgtagcaaacacgacaccgatataatctggaacaagtatccagcctacgcgttcacctcacctacgagggcc

Figure 16 (cont.)

agccgacaatattctaccgcgactacgaggagtggtcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccggagga  
agcactgacatcgttactacgacaacgacgagctgatatctgtgagaacggctacggaagcaagccgggactgataacatacatcaacctc  
gcctcaagcggaagccggaaggtgggtctacgttccgaagttcgccggagcgtgcatccacgagtagacacggcaacctcgccggctgggtgg  
acaagtggtgggtggaagcggtgggtgtacctgaggccctgcccacgacccggccaacggctattacggctactccgtctggagctatt  
gcggtgtgggtga

SEQ ID NO.: 4

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Glu Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 5

atggccaagtactccgagctggaagagggcggttataatcgaggccttacttgggacgtcccagggtggaggaatctggtgggacaccat  
caggagcaagataccgagtggtacgagggcggaatatccgccatttggattccccggcaagcaagggcatggcgccctattcgcgatg  
ggctacgaccctacgacttctttgacctcgggtgagtagcaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacaccgcccacgcctatggcatgaaggtatagccgatatagtcatcaaccaccgcgccggcggtgacctggagtgaacccct  
cgtgaacgactatactggaccgacttctcaaaggtcgcgtcgggtaatacacggccaactacctgacttccaccgaacgagctccatgc  
ggcgattccgggaacatttggaggctatcccgacatatgccacgacaagagctgggaccagtactggctctggccagccaggagctac  
gcggcatatctcaggagcatcggcatcgcgtgctgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactggt  
ggggaggctggcggttgagagtactgggacaccaacgtcgcgtgttctcaactgggcatactcgagcgggtccaaggtctttgacttcg  
ccctactacaagatggatgaggcctttgacaacaaaacattccagcgcgtcgtcttgccttcagaacggccaagctgttctccgcgac  
ccgtcaaggccgtaaccttttagcaaacacgacaccgatataatctggaacaagtaccttgccttatgccttacctacctaagggccag  
ccgcatattctaccgcgaccacgaggagtggctcaacaaggacaggtgaacaacctcatatggatacacgaccacctcgaggtggaag  
caccgacatagctactacgataacgatgaactcatcttcaggaacgggtacggggacaagccggggctataacctacatcaacctaggc  
tcgagcaaggccggaaggtgggttatgtgccgaagttcgggcgctgcatccacgagtagtactggtaacctcgaggtgggttagacaa  
gtacgtctactcaagcggctgggtctatctcgaagctccagcttacgacctgccaacgggcagtagtggtactccgtgtggagctactgcggg  
gtgggctga

SEQ ID NO.: 6

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly

Figure 16 (cont.)

Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Leu Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp His Glu Glu Trp Leu Asn  
Lys Asp Arg Leu Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 9

atggccaagtactccgagctggaagagggcgggctcataatgcaggccttctactgggacgtcccatgggaggaatctggtgggacacgat  
agcccagaagatacccgaactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcatggcgggcgccctattcgatg  
ggctacgaccctacgacttcttgacctgggtgagtacgaccagaagggaaacggtagagacgcgctttggctccaagcaggagctctgtgaa  
catgataaacacggcccatgctacggcataaaggtcatagcggacatcgtcataaaccaccgcgcaggcggagacctcgaagtgaacccg  
ttcgttggggactacacctggacggacttctcaaggtggcctcgggcaatatactgccaactacctgacttccacccgaacgagctccatg  
cggcgcatcgcgaacatttgaggctatcccgacatatgccacgacaagagctgggaccagtactggctctgggcccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgtgcctggcgcttcgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactggt  
ggggaggctggcggttgagagtactgggacaccaacgtcgacgctgttctcaactgggcatactcgagcgggtgccaaggtcttgacttcg  
cccttactacaagatggacgagccctcgataacaacaacattccccgccctgggtggacgccctcagatacggtcagacagtggtcagccgcg  
accggtcaaggtgtgacgtttgtagccaaccacgataccgataatactggaacaagtatccagctacgcgttcactctcacctacgagggc  
cagccgacaatattctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccggagg  
aagcactgacatcgtttactacgacaacgacgagctgatattcgcgagaaacggctacggaagcaagccgggactgataacatacatcaacct  
cgccctcaagcaagccggaaggtgggtttacgttcgaagttcgaggctcgtgcatacacgagtacaccggcaatctcggcggtgggtgg  
acaagtggtgggactcaagcgggtgggtctacctcgaggctcctgccacgaccggccaacggccagtacggctactccgtctggagctac  
tgcggtgttggtga

SEQ ID NO.: 10

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Ala Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu

Figure 16 (cont.)

Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 11

atggccaagtacctggagctcaggagggcgggctcataatgcaggccttctactgggacgtcccatgggaggaatctggtgggacacgat  
agcccagaagatacccgactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcatgggcggcgctattcgtatg  
ggctacgaccttacgacttcttgacctcggtagtacgaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacaccgccacgctatggcatgaaggtaatagccgatatagtcatcaaccaccgcgccggcggtgacctggagtggaaacctt  
cgtgaacgactatacctggaccgacttctcaaaggctcgcgtcgggtaatacacggccaactacctcgaactccaccgaacgagctccatgc  
ggcgattccgggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggccaggcaggagactac  
gcggcatatctcaggagcatcgccatcgatgctggcgcttcgactacgtcaagggtatgtccctgggtcgtcaaggactggctgaactggt  
ggggaggctgggcgggtggagagtactgggacaccaacgtcgacgtgtttcctaactgggcatactcagcgggtccaaggtctttgacttcg  
cccttactacaagatggacgaggccttcgataacaacaacattccccgcctggtggacgccctcagatacggtcagacagtgtgcagccgcg  
accggtcaaggctgtgacgtttgtgccaaccacgataccgataatactggaacaagtatccagcctacgcgttcacctcacctacgagggc  
cagccgacaatatttaccgcgactacgaggagtggctcaacaaggatacgtcaagaacctcatctggatacatgacaacctgccggagg  
aagcacgagcatagttactacgacagcgacgagatgatcttcgtgaggaacggctatggaagcaagcctggccttataacttacatcaacctc  
ggctcagcaaggttgaaggtgggtctacgttccgaagttcgcgggagcgtgcatccacgagtacaccggcaacctcgcggtggtgg  
acaagtgggtgactcaagcggtgggtgtacctcgaggccctgccacgacccggccaacggctattacggctactccgtctggagctac  
tgcggtgttgctga

SEQ ID NO.: 12

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Thr Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Ser Ile Val Tyr  
Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 13

atggccaagtacctggagctcgaagagggcggggtcataatgcaggcgttctactgggacgtgccttcaggaggaatatggtgggacacaat  
acggcagaagataccggagtgtacgatgccggaatctccgcaatatggattccccggcgagcaagggcatgggcggcgctattcgtatg  
ggctacgaccttacgacttcttgacctcggtagtatgaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
atgataaacacggcacatgcctacggcataaaggctcatagcgacatcgtcataaaccaccgcgcaggcggagacctcgagtgaacctgtt  
cgttggggactacacctggacggacttctcaaagggtggcctcgggcaatatactgccaactacctcgaacttccacccaacgaggtcaagt  
ctgtgacgagggcacatttggaggcttccagacatagccacgagaagagctgggaccagcactggctctgggcgagcgtgagagctac  
gccgctactctaaggagcatcggcgttgatgcctggcgcttcgactacgtcaagggtacggagcgtgggtcgtcaaggactggctggactg  
gtggggaggctgggcgctcggggagtactgggacacaaacgttgatgcactgtcactgggcctactcgagcgtgcaaaagtcttcgactt

Figure 16 (cont.)

cccgtctactacaagatggatgagcccttgacaacaaaacattccagcgctcgtctctgcccttcagaacggccagactgttgtctcccgcg  
accggttcaaggccgtaacctttagcaaacacgacaccgatataatctggaacaagtatccagcctacgcgttcacctacacgaggg  
ccagccgacaatatctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgcggag  
gaagcactgacatagtctactacgataacgatgaactcatcttcgtaggaacgggtacggggacaagccggggcttataacctacatcaacct  
aggctcgaagcaggccggaagggtgggttatgtgccgaagttcgcgggcgctgcacccagagtatactggaacctcggagggtgggttag  
acaagtacgtctactcaagcggctgggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgtgtggagctactg  
cgggttggctga

SEQ ID NO.: 14

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp  
Val Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp  
Glu Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val  
Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 15

atggccaagtactccgagctggaagagggcgggctcataatcgaggccttactgggacgtccccatgggaggaatctggtgggacacgat  
agcccagaagatacccgactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcagtgggcgccctattcagatg  
ggctacgacctacgacttctttgacctcgggtgagtagcaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaggatcagcggacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccg  
ttcgttggggactacacctggacggacttctcaaagggtggcctcgggcaaatatactgccaaactacctcgaactccaccgaaacgagctccatg  
cgggcgattccggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgtgcctggcgcttcgactacgtcaagggtacggagcgtgggtcgtcaaggactggctggactg  
gtggggagggtgggcccgtcggggagtagtgggacacaaacgttgatgcactgctcaactgggcctactcgagcgtgcaaaagcttcgactt  
cccgtctactacaagatggatgagcccttgacaacaaaacattccagcgctcgtctctgcccttcagaacggccagactgttctcccgcg  
accggttcaaggccgtaacctttagcaaacacgacaccgatataattggaacaagtaccggcctacgccttcacctacacacgaggg  
ccagccgacgatattctaccgcgactacgaggagtggctcaacaaggacaggctcaagaacctcatctggatacacgaccacctgcccgtg  
gaagcactgacatcgttactacgacaacgacgagctgatattcgtgagaacggctacggaagcaagccgggactgataacatacatcaacc  
tcgcctcaagcaaaagccggaagggtgggttatgtgccgaagttcgcgggcgctgcacccagagtatactggaacctcggagggtgggttag  
acaagtacgtctactcaagcggctgggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgtgtggagctattgc  
ggtgttgggtga

SEQ ID NO.: 16

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp

Figure 16 (cont.)

Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val  
Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Arg Leu Lys Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 17

atggccaagtactccgagctggaagggggcgggctcataatgcaggccttactgggacgtccccatgggaggaatctggtgggacacgat  
agccagaagatacccgactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcatgggcggcgcttattcgatg  
ggctacgaccctacgacttctttgacctcggtgagtagcaccaggagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaggctcatagcggacatcgtcataaaccaccgcgcaggcggagacctcgagtgaacccg  
ttcgttggggactacacctggacggacttctcaaaggtggcctcgggcaaatatactgccaaactacctcgacttccacccaacgagggtcaagt  
gctgtgacgagggcacatttggaggcttcccagacatagcccacgagaagagctgggaccagcactggctctgggagcagcagatgagagcta  
cgccgctacctaaggagcatcgccgttgatgcctggcgcttcgactacgtcaagggctacggagcgtgggtcgtcaaggactggctggact  
ggtggggaggtgggcccgtcggggagtactgggacacaaacgttgatgcactgctcaactgggcctactcgagcgatgcaaaagtcttcgac  
ttcccgctactacaagatggacgcggcctttgacaacaagaacattccgcactcgtcgaggccctcaagaacgggggcacagtcgtcagc  
cgcgacccgtttaaggccgtaaccttcgttgcaaacacgacaccgatataatctggaacaagtatccagcctacgcgttcacctcacctacga  
gggcccagccgacaatattctaccgcgactacgaggagtgggtcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccg  
gaggaagcacgagcatagttactacgacagcgagatgattcgtgaggaacggctatggaagcaagcctggccttataacttacatcaa  
cctcggtcagcaaggttgaaggtgggtttacgtccgaagttcgcaggctcgtgcatacacgagtacaccggcaatctcggcggtgggt  
ggacaagtgggtggactcaagcggtgggtctacctcgaggctcctgccacgaccggccaacggccagtagcggtactccgctctggagc  
tactcggtgttgggtga

SEQ ID NO.: 18

Met Ala Lys Tyr Ser Glu Leu Glu Gly Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Glu Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp  
Val Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp  
Ala Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Ser Ile Val

Figure 16 (cont.)

Tyr Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 19

atggccaagtacctggagctcgaagagggcgggctcataatgcaggccttctactgggacgtcccatgggaggaatctggtgggacacgat  
agcccagaagatacccgactgggcaagcgccgggatttcggcgatatggattcctcccgcgagcaagggtatgagcggcggtattcgatgg  
gctacgacctctacgattattttgaccttggtgagtactaccagaagggaacgggtgaaacgaggttcggctcaaagcaggagctcataaacat  
gataaacacggcccatgcctacggcataaaggctatagcggacatcgtcataaaccaccgcgcaggcggagacctcgagtgaacccgttc  
gttggggactacacctggacggacttctcaaagggtggcctcgggcaatatactgccaactacctcgacttccacccgaacgagctccatgcg  
ggcgattccggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctggggccagccaggagagctacgc  
ggcatatctcaggagcatcggtcgtgctgctgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactggtgg  
gggggctggcggttgagagtactgggacaccaacgtcgacgtgttctcaactgggcatactcgagcgggtccaaggctcttgacttgcgc  
ctctactacaagatgtaggagccttgacaacaaaaacattccagcgtcgtctctgcccctcagaacggccagactgttctctccgcgaccc  
gttcaaggccgtaaccttttagcaaacacacgacaccgatataatttgaacaagtaccggcctacgcttcatctcacctacgaggggccag  
ccgacgatattctaccgcgactacgaggagtggctcaacaaggacaggtcaagaacctcatctggatacacgaccacctcgccggtggaag  
cactgacatcgtttactacgacaacgacgagctgatattcgtgagaacggctacggaagcaagccgggactgataacatacatcaacctcgc  
ctcaagcaaagccggaaggtgggttatgtgccgaagttcgcgggcgctgcatccacgagcatactggttaacctcgagggtgggtagaca  
agtacgtctactcaagcggctgggtctatctcgaagctccagcttaccacctgccaacgggcagtatggctactccgtgtggagctactcggg  
tgttggctga

SEQ ID NO.: 20

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val  
Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu Ala  
Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg Asp  
Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr  
Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Arg Leu Lys Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr  
Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu His  
Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro  
Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 21

atggccaagtactccgagctggaagagggcgggcgttataatgcaggccttctactgggacgtcccaggtggaggaatctggtgggacacat  
caggagcaagataccggagtggtagaggcggggaatatccgccatttggattcctccgggagcaagggtatgagcggcggtattcgatgg  
gctacgacctctacgatatttggacctgggtgagtactaccagaagggaacgggtgaaacgaggttcggctcaaagcaggagctcataaac  
atgataaacacggcccatgcctacggcataaaggctatagcggacatcgtcataaaccaccgcgcaggcggagacctcgagtgaacccgtt  
cgttggggactacacctggacggacttctcaaagggtggcctcgggcaatatactgccaactacctcgacttccacccgaacgagctccatgc  
ggcgattccggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctggggccagccaggagagctac

Figure 16 (cont.)

gcggtatatctcaggagcatcggcatcgatgcctggcgcttcgactacgtcaagggtacggagcgtgggtcgtcaaggactggctggactg  
gtggggaggctgggcccgtcggggagtactgggacacaaacgttgatgcactgctcaactgggcctactcgagcgtgcaaaagtcttcgactt  
cccgtctactacaagatggatgaggcctttgacaacaaaaacattccagcgctcgtctctgcccttcagaacggccagactgttctccccgcg  
accgttcaaggccgtaaccttttagcaaacacgacaccgatataatttggaacaagtagccggcctacgccttcacctacacgaggg  
ccagccgacgatattctaccgagactacgaggagtggctcaacaaggacaggctcaagaacctcatctggatacacgactacctcgccggtg  
gaagcactgacatcggttactacgacaacgacgagctgatattcgtgagaacggctacggaagcaagccgggactgataacatacatcaacc  
tcgctcaagcaaaagccggaaggtgggttatgtgccgaagttcgcgggcgctgcatccacgagtatactggtaacctcgagggctgggtg  
acaagtacgtctactcaagcggtgggtctatctcgaagctccagcttacgacctgccaacgggcagtagtggtactccgtgtggagctattgc  
gggttggctga

SEQ ID NO.: 22

Met Ala Lys Tyr Ser Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Gly Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Asp Leu Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Val  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val  
Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Arg Leu Lys Asn Leu Ile Trp Ile His Asp Tyr Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 23

atggccaagtactccgagctggaagagggcggcgttatagtgcaggccttactgggacgtcccaggtggaggaatctgggtgggacaccat  
caggagcaagataccggagtgttacgaggcgggaatatccgccatttgattccccggcgagcaagggcagtgggcggcgcctattcgatg  
ggctacgacctacgacttctttgacctcggtagtacgaccagaagggaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgacctacggcataaaggtcatagcgacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccg  
ttcgttggggactacacctggacggacttctcaaaaggtggcctcgggcaataatactgccaactacctcgaactccacccgaacgagctccatg  
cgggcgattccggaacatttgaggctatcccgacatatgccacgacaagagctgggaccagtagtggctctgggccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgatgcctggcgcttcgactacgtcaagggtacggagcgtgggtcgtcaaggactggctggactg  
gtggggaggctgggcccgtcggggagtactgggacacaaacgttgatgcactgctcaactgggcctactcgagcgtgcaaaagtcttcgactt  
cccgtctactacaagatggatgaggcctttgacaacaaaaacattccagcgctcgtctctgcccttcagaacggccagactgttctccccgcg  
accgttcaaggccgtaaccttttagcaaacacgacaccgatataatctggaacaagtatccagcctacgcgttcacctcacctacgaggg  
ccagccgacaatatctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccggag  
gaagcatgagcatagttactacgacagcgacgagatgatcttcgtgaggaacggctatggaagcaagcctggccttataactatcatcaacct  
ggctcgagcaaggttggaaggtgggtctacgttccgaagttcgcgggagcgtgcatccacgagtacaccggcaacctcggcggtgggtg  
acaagtggtggactcaagcgggtgggtgtacctcgaggccccctgccacgacccggccaacggctattacggctactccgtctggagctatt  
gcggtgttggtga

SEQ ID NO.: 24

Figure 16 (cont.)

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Val Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val  
Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Met Ser Ile Val Tyr  
Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 25

atggccaagtacctggagctcgaagagggcgggctcataatgcaggccttctactgggacgtcccatgggaggaatctgggtgggacacgat  
agcccagaagatacccgactgggcaagcgccgggatttcggcgatatggattcctcccgcgagcaagggtatgagcggcggtattcgatgg  
gctacgacccctacgattattttgacctcggtgagtactaccagaagggaacgggtgaaacgaggttcggctcaaagcaggagctcataaacat  
gataaacaccgcccacgcctatggcatgaaggtatagccgatatagtcatcaaccaccgcgccggcggtgacctggagtgaacccctcgt  
gaacgactatacctggaccgacttctcaaaggctcgcgtcgggtaatacacggccaactacctgacttccaccggaacgagctccatcgggg  
cgattccgggaacatttggaggctatcccgcataatgccacgacaagagctgggaccagtactggctctgggccagccaggagactacgcgg  
catatctcaggagcatcggcatcgtgcctggcgcttcgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactgggtggg  
aggctgggcggttggagagtactgggacaccaacgtcgacgctgttctcaactgggcatactcgagcgggtccaagggtcttgacttcgccctc  
tactacaagatggacgaggccttcgataacaacaacattcccgccttgggtgggcccctcagatacggtcagacagtggtcagccgcgaccc  
gttcaaggctgtgacgtttgtagccaaccacgataccgatataatctggaacaagatccagcctacgcgttcacctcacctacgagggccagc  
cgacaatattctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgcggaggaagc  
accgacatagtctactacgataacgatgaactcatcttcgtaggcacggctacggggacaagccggggcttataacctacatcaacctaggct  
cgagcaaggccggaagggtgggttacgttccgaagttcgaggctcgtgcatacacgagtacaccggcaatctcggcgggtgggtgggaca  
gtgggtggactcaagcgggtgggtctacctcgaggctcctgccacgacccggccaacggccagtacggctactccgtctggagctattgca  
gtgttgggtga

SEQ ID NO.: 26

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val  
Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu Ala  
Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Gly Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp

Figure 16 (cont.)

Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr  
Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr  
Asp Asn Asp Glu Leu Ile Phe Val Arg His Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr  
Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro  
Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 27

atggcaaatgattccgagctcgaagagggcggttataatgcaggccttctactgggacgtccaggtggaggaatctggtgggacaccatc  
aggagcaagataccggagtgtgtacgagggcggaataatccgccatttgattctctccgcgagcaaggggtatgagcgggcggtattcgatggg  
ctacgaccctacgattatgttgacctcggtgagtactaccagaaggggaacgggtggaacgaggttcgggtcaaagcaggagctcataaacatg  
ataaacacggcccatgctacggcataaaggtcatagcggacatcgtcataaaccaccgcgcaggcggagacctcgaagtgaacccgttcgt  
tggggactacacctggagcgacttctcaaggtggcctcgggcaatatactgccaactacctcgacttccaccgaacgagctccatcgggg  
cgattccggaacatttgaggctatcccgcacatgccacgacaagagctgggaccagtactggctctgggcccagccaggagagctacgcgg  
catatctcaggagcatcggcatcgatgcctggcgcttcgactacgtcaagggctatgtccctgggtcgtcaaggactggctgaactggtggg  
aggctgggcggttgagagtagtactgggacaccaacgtcgacgtgttctcaactgggcatactcagcggtgccaaggtcttgacttcgcctc  
tactacaagatggagcggcgcttgacaacaagaacattcccgactcgtcgaggccctcaagaacgggggcacagtcgtcagccgcgacc  
cgtttaaggccgtaaccttcgttgcaaacacgacaccgatataatctggaacaagatccagcctacgcgttcacctacacagggccag  
ccgacaatattctaccgcgactacgaggagtggctcaacaagataagctcaagaacctcatctggatacatgacaacctcgccggaggaag  
cactgacatcgtttactacgacaacgacgagctgatattcgtgaaacgggtacggaagcaagccgggactgataacatacatcaacctcgc  
gtcaagcaaagccggaaggtgggttacgttcgaagttcgaggctcgtgcatacacgagtagaccggcaatctcggcggtgggtggaca  
agtgggtggactcaagcgggtgggtctacctcgaggctcgtcccgacccggccaacggccagtaggctactccgtctggagctactgc  
ggtgttgggtga

SEQ ID NO.: 28

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val  
Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Ala Ala  
Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser Arg Asp  
Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr  
Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr  
Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr  
Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro  
Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 29

atggccaagtacctggagctcgaagagggcggttcataatgcaggccttctactgggacgtcccatgggaggaatctggtgggacacgg  
agcccaagaagataccgactgggcaagcgccgggatttcggcgatatggattccccggcgagcaagggcatgggcggcgcttattcgatg  
ggctacgaccctacgacttcttgacctcggtgagtacgaccagaaggggaacggtagagacgcgtttggctccaagcaggagctcgtgaa

Figure 16 (cont.)

catgataaacacggcccatgcctacggcataaaaggtcatagcggacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccg  
ttcgttggggactacacctggacggacttctcaaaggttggtctcgggcaaatatactgccaactacctcgacttccacccgaacgagtcctatgc  
ggcgattccggaacatttggaggctatcccacacgacaagagctgggaccagtactggctctgggccagccaggagagctac  
gcggcataatctcaggagcatcggcatcgcctggcgttcgactacgtcaagggtatgtccctgggtcgtcaaggactggctgaactggt  
ggggaggctgggctgtggagagtactgggacaccaacgtcgacgctgttctcaactgggcatactcagcgggtccaaggctttgacttcg  
cccttactacaagatggatgaggcctttgacaacaaaacattccagcgtcgtctcgtcccttcagaacggccagactgttgcctcccgac  
ccgttcaaggccgtaacctttgtagcaaacacgacaccgatataatctggaacaagtaccttgccttatgccttcacctcacctacgaaggccag  
cccgcatattctaccgcgactacgaggagtggctcaacaaggacaggttgaacaacctcatatggatacacgaccacctcgcagggggaag  
caccgacatagctactacgataacgatgaactcatcttcgtaggaacggctacggggacaagccggggttataacctacatcaacctaggc  
tcgagcaaggccggaagggtgggtttatgtgccgaagttcgcggcgcggtcatccacgagtatactggttaacctcgagggtggtagacaa  
gtacgtctactcaagcggctgggtctatctcgaagctccagcttacgaccttgccaacgggcagtatggctactccgtgtggagctactcggt  
gttgggtga

SEQ ID NO.: 30

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met  
Gly Gly Ile Trp Trp Asp Thr Val Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Val Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Leu Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Arg Leu Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 31

atggcaagtagtccgagctggaagagggcggcgttataatgcaggccttctactgggacgtcccagggtggaggaatctggtgggacacat  
caggagcaggataccggagtggtacgagggcgggaatatccgccatttgattccccggcgagcaagggcattggcgccgctattcgatg  
ggctacgaccctacgacttcttgacctcggtagtagtacgaccagaagggaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaaggtcatagcggacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccg  
ttcgttggggactacacctggacggacttctcaaaggttgccctcgggcaaatatactgccaactacctcgacttccacccgaacgagtcctatg  
cggcgattccggaacatttggaggctatcccacacgacaagagctgggaccagtactggctctgggccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgcctggcgtttgactacgtgaagggtacggagcgtgggtcgtcaaggactggctcaactgg  
tggggcggttggcggttggcgagtactgggacaccaacgttgatgcactcctcaactgggcctactcagcggcgccaaggcttctgacttc  
ccgcttactacaagatggacgagggccttcgataacaacaattccgcctgggtggacgccctcagatacggtcagacagtgggtcagccgc  
gaccggtcaaggctgtgacgtttgtagccaaccacgataccgatataatctggaacaagtatccagcctacgcgttcatctcacctacgaggg  
ccagccgacaatttctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctggccggag  
gaagcacgagcatagtttactacgacagcgacgagatgatcttctgtaggaccggctatggaagcaagcctggccttataacttacatcaacct  
cggctcagcaagggttgaagggtgggtttatgtgccgaagttcgcggcgcggtcatccacgagtatactggttaacctcgagggtgggtaga  
caagtacgtctactcaagcgggtggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgtgtggagctattgcg  
gttgggtga

Figure 16 (cont.)

SEQ ID NO.: 32

Met Ala Lys Tyr Ser Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Arg Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Ser Ile Val Tyr  
Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Thr Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 33

atggccaagtactccgagctggaagagggcggggtcataatgcaggcggttctactgggacgtgccttcaggaggaatatggtgggacacaat  
acggcagaagataccggagtgtacgatgccggaatctccgcaatatggattcctcccgagcaagggtatgagcggcggtattcgtatgg  
gctacgacccctacgattatttgcctcgggtgagtactaccagaaggggaacgggtgaaacgaggttcggctcaaagcaggagctcataaacat  
gataaacacggcccatgcctacggcataaaggatcatagcggacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccgttc  
gttggggactacacctggacggacttctcaaaggtggcctcgggcaaatatactgccaactacctcgacttcacccgaacgagctccatgcg  
ggcgattccgggaacatttggaggctatcccgacatagccacgacaagagctgggaccagtactggctctgggcccagccaggagctacgc  
ggcatatctcaggagcatcggcatcgtatgcctggcgcttgactacgtgaagggtctacggagcgtgggtcgtcaaggactggctcaactgggtg  
ggcggtggtggcggttggtgagctactgggacaccaacgttgatgcactcctcaactgggctactcgagcggcgccaaggtcttcgactttcc  
gcttactacaagatggacgcggcctttgacaacaagaacattcccgactcgtcaggccctcaagaacgggggcacagtcgtcagccgcg  
accggttaaggccgtaaccttcgttgcacacacgacaccgatataatctggaccaagtaccttgcttatgctttcatcctcacctacgaaggcca  
gcccgtcatattctaccgcgactacgaggagtgggtcaacaaggacaggttgaacaacctcatatggatacacgaccacctcgcaggtggaag  
caccgacatagtctactacgataacgatgaactcatctcgtcaggaaacggctacggggacaagccggggcttataacctacatcaacctaggc  
tcgagcaaggccggaaggtgggtttacgttccgaagttcgcaggctcgtgcatacacgagtagacaccggcaatctcggcggtgggtgggaca  
gtgggtggactcaagcgggtgggtctacctcgaagctcctgcccacgacccggccaacggccagtagcgtactccgtctggagctactgcg  
gtgttgctga

SEQ ID NO.: 34

Met Ala Lys Tyr Ser Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala

Figure 16 (cont.)

Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Ala  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Thr Lys Tyr Leu Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Arg Leu Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 35

atggccaagtactccgagctggaagagggcggtataatgcaggccttactgggacgtcccaggtggaggaatctggtgggacaccat  
caggagcaagataccggagtggtacgagggggaatatccgccatttgattccccggcgagcaagggcatgggcggcctattcgatg  
ggctacgacccctacgacttcttgacctcggtgagtagcaccagaagggaacggtagagacgcgcttggctccaagcaggagctcgtgaa  
catgataaacaccgcccacgcctacggcatcaaggtcatcgacacatagtaataaccaccgcgccggaggagacctgagtgaacccct  
tcgtcaatgactacacctggacggactctcgaaggtcgcttcggcaagtagacacggccaactacctcgactccacccaacagggtcaagtg  
ctgtgacgagggcacatttgagggttcccagacatagcccaagagaagagctgggaccagcactgggtctggcgagcgatgagagctac  
gccgctacctaaggagcatcgcggtgatgcctggcgcttcgactacgtcaagggtctgctcctgggtcgtaaggactggctgaactggt  
ggggagggtggcggttgagtagtactgggacaccaacgtcgacgctgttctcaactgggcatactgagcggtgccaaggcttctgactcg  
ccctactacaagatggacgcggccttgacaacaagaacattccgcactcgtcgaggccctcaagaacgggggcacagtcgtcagccgc  
gaccggttaaggccgtaaccttcgttgcaaacacgacacgatataatctggaacaagtatccagcctacgcgttcacctacacgaggg  
ccagccgacaatattctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctatctggatacatgacaacgtcgccggag  
gaagcaccgacatagctactacgataacgatgaactcatcttcgcaggaaacggctacggggacaagccggggcttataacctacatcaacct  
aggctcgagcaaggccggaagggtgggtttacgttccgaagttcgaggctcgtgcatacacgagtagacaccggcaatctcgccggctgggtgg  
acaagtggtggactcaagcggtgggtctacctcgaggctcctgccacgaccggccaacggccagtagcgtactccgtctggagctac  
tgcggtgtgggtga

SEQ ID NO.: 36

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp  
Val Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp  
Ala Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Val Ala Gly Gly Ser Thr Asp Ile Val  
Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 71

atggccaagtactccgagctcgaagagggcggtcataatgcaggcgttactgggacgtgccttcaggaggaatatggtgggacacaat

Figure 16 (cont.)

acggcagaagataccggagtggtagatgccggaatctccgaatatggattccccggcgagcaagggcatgggcggcgcctattcga  
ggctacgaccttacgacttctttgacctcggtagtagtagaccagaaggggaacggtagagacgcctttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaggtagtagcggacatcgtcataaaccaccgcgcaggcggagacctcgaaggaaacccg  
ttcgttggggactacacctggacggacttctcaaaggtagcctcgggcaaatatactgccaactacctcgaactccacccgaacgagctccatg  
cgggctgattccggaacatttggaggctatcccgacatatgccacgacaagagctgggaccagtactggctctgggccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgaatcggcgttcgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactggt  
ggggaggctggggtgttgagagtagtgggacaccaacgtcgcgtgttctcaactgggcatactcgaagggtccaaggctttgacttcg  
cccttactacaagatggatgaggcctttgacaacaaaaacattccagcgtcgtctcgtcccttcagaacggccagactgtgtctcccgac  
ccgtcaaggccgtaacctttgtagcaaacaccgacaccgatataatctggaacaagtatccagcctacgcgttcacctacgagggcc  
agccgacaataattctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcggcgagga  
agcactgacatcgtttactacgacaacgacgagctgatattcgtgagaacgggtacggaagcaagccgggactgataacatacatcaacctc  
gcctcaagcaaagccggaagggtgggtttatgtgccgaagttcggggcggtgcatccacgagataactgtaacctcggaggctgggtaga  
caagtacgttactcaagcggctgggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgtgtgagctactgc  
gggtgggtgta

SEQ ID NO.: 72

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Val Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 49

gtggtttatgacgatgtccgctatgacctttatgccgtaggcatgggcccgtgtttatcatgttcacgagctcctgcttgagccaaagcgcgtctc  
accgttcccttctggtcgtactcaccgagggtcaaagaagtcgtaggggtcgtagcccatcgaataggcggcccatgcccttgctcgccggg  
ggaatccatatcgccgaaatcccggtcgtgccagtcgggtatcttctgggctatcgtgtcccaccagattcctcccatggggacgtccagta  
gaaggcctgcattatgagcccgcctcttcgagcccgaatactttgccataagttacctctactagtagattaaaattctgttctgtgtgaaatt  
gtt

SEQ ID NO.: 50

Val Val Tyr Asp Asp Val Arg Tyr Asp Leu Tyr Ala Val Gly Met Gly Arg Val Tyr His Val His Glu  
Leu Leu Leu Gly Ala Lys Ala Arg Leu Tyr Arg Ser Leu Leu Val Val Leu Thr Glu Val Lys Glu Val  
Val Gly Val Val Ala His Arg Ile Gly Ala Ala His Ala Leu Ala Arg Arg Gly Asn Pro Tyr Arg Arg  
Asn Pro Gly Ala Cys Pro Val Gly Tyr Leu Leu Gly Tyr Arg Val Pro Pro Asp Ser Ser His Gly Asp  
Val Pro Val Glu Gly Leu His Tyr Glu Pro Ala Leu Phe Glu Pro Gly Ile Leu Cys His Lys Leu Pro  
Pro Thr Ser Arg Leu Lys Phe Cys Phe Leu Cys Glu Ile Val

Figure 16 (cont.)

SEQ ID NO.: 51

ATGGCCAAGTACCTGGAGCTCGAAGAGGGCGGGGTCATAATGCAGGCGTTCTACTGGG  
ACGTGCCTTCAGGAGGAATATGGTGGGACACAATACGGCAGAAGATACCGGAGTGGT  
ACGATGCCGGAATCTCCGCAATATGGATTCCCCCGGCGAGCAAGGGCATGGGCGGCGC  
CTATTCGATGGGCTACGACCCCTACGACTTCTTTGACCTCGGTGAGTACGACCAGAAG  
GGAACGGTAGAGACGCGCTTTGGCTCCAAGCAGGAGCTCGTGAACATGATAAACACC  
GCCCACGCCTATGGCATGAAGGTAATAGCCGATATAGTCATCAACCACCGCGCCGGCG  
GTGACCTGGAGTGGAACCCCTTCGTGAACGACTATACCTGGACCGACTTCTCAAAGGT  
CGCGTCGGGTAAATACACGGCCAACCTACCTCGACTTCCACCCCAACGAGGTCAAGTGC  
TGTGACGAGGGCACATTTGGAGGCTTCCAGACATAGCCCACGAGAAGAGCTGGGAC  
CAGCACTGGCTCTGGGCGAGCGATGAGAGCTACGCCGCCTACCTAAGGAGCATCGGCG  
TTGATGCCTGGCGCTTTGACTACGTGAAGGGCTACGGAGCGTGGGTCTGCAAGGACTG  
GCTCAACTGGTGGGGCGGCTGGGCCGTTGGCGAGTACTGGGACACCAACGTTGATGCA  
CTCCTCAACTGGGCCTACTCGAGCGGCGCCAAGGTCTTCGACTTCCCGCTCTACTACAA  
GATGGATGAGGCCTTTGACAACAAAAACATTCCAGCGCTCGTCTCTGCCCTCAGAAC  
GGCCAGACTGTTGTCTCCCGCGACCCGTTCAAGGCCGTAACCTTTGTAGCAAACCACG  
ACACCGATATAATCTGGAACAAGTATCCAGCCTACGCGTTCATCCTCACCTACGAGGG  
CCAGCCGACAATATTCTACCGCGACTACGAGGAGTGGCTCAACAAGGATAAGCTCAAG  
AACCTCATCTGGATACATGACAACCTCGCCGGAGGAAGCACTGACATCGTTTACTACG  
ACAACGACGAGCTGATATTCGTGAGAAACGGCTACGGAAGCAAGCCGGGACTGATAA  
CATACTCAACCTCGCCTCAAGCAAAGCCGGAAGGTGGGTTTACGTTCCGAAGTTCGC  
AGGCTCGTGCATACACGAGTACACCGGCAATCTCGGCGGCTGGGTGGACAAGTGGGTG  
GACTCAAGCGGCTGGGTCTACCTCGAGGCTCCTGCCACGACCCGGCCAACGGCCAGT  
ACGGCTACTCCGTCTGGAGCTATTGCGGTGTTGGCTGA

SEQ ID NO.: 52

MAKYLELEEGGVIMQAFYWDVPSGGIWWDTIRQKIPEWYDAGISAIWIPPASKGMGGAYS  
MGYDPYDFFDLGEYDQKGTVETRFSGSKQELVNMINTAHAYGMKVIADIVINHRAGGDLE  
WNPVNDYTWTDFSKVASGKYTANYLDFHPNEVKCCDEGTFGGFPDIAHEKSWDQHWL  
WASDESYAAYLRSIGVDAWRFDYVKGYGAWVVKDWLNWWGGWAVGEYWDTNVDAL  
LNWAYSSGAKVFDPLYYKMDEAFDNKNIPALVSALQNGQTVVSRDPFKA VTFVANHDT  
DIIWNKYPAYAFILTYEQPTIFYRDYEEWLNKDKLKNLIWIHDNLGGSTDIVYYDNDELI  
FVRNGYGSKPGLITYINLASSKAGRWVYVPKFAGSCIHEYTGNLGGWVDKWVDSSGWVY  
LEAPAHDPANGQYGYSVWSYCGVG

SEQ ID NO.: 37

atggccaagtacctggagctcgaagagggcggggtcataatgcaggcgttactgggacgtgccttcaggaggaatatggtgggacacaat  
acggcagaagataccggagtggtagatgccggaatctccgcaatatggattccccggcgagcaagggcatggcggcgcctattcgatg  
ggctacgaccctacgacttctttgacctcggtagtagtagcagaccagaagggaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacaccgcccacgcctatggcatgaaggtaatagccgatatagtcatcaaccaccgcgcggcggtgacctggagtgaacccctt  
cgtgaacgactatacctggaccgacttctcaaaggtcgcgtcgggtaatacacggccaactacctcacttccaccgaacgagctccatgc  
ggcgattccggaacatttgaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggcccagccaggagagctac  
gcggcatatctcaggagcatcggcatcgtgctggcgctttgactacgtgaagggtacggagcgcgggtcgtcaaggactggctcaactg  
gtggggcggtgggcccgttgccgagtagtgggacaccaacgttgatgcactcctcaactgggctactcgagcggcgccaaggtcttcgactt  
cccgtctactacaagatggatgagggcctttgacaacaaaaacattccagcgctcgtctctgccctcagaacggccagactgttctcccgcg  
accgttcaaggccgtaaccttttagcaaacaccgacaccgatataatctggaacaagtatccagcctacgcgttcactcctacctacgaggg  
ccagccgacaatattctatcgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccggagg  
aagcactgacatcgttactacgacaacgacgagctgatattcgtgagaacggctacggaagcaagccgggactgataacatacatcaact  
cgctcaagcaaagccggaaggtgggttacgttccgaagttcgcaggctcgtgcatacacgagtacaccggcaatctcgccggctgggtg

Figure 16 (cont.)

acaagtgggtggactcaagcggctgggtctacctgaggctcctgccacgacccggccaacggccagctacggctactccgtctggagctac  
tgcgggggtgggtga

SEQ ID NO.: 38

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly  
Tyr Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Arg Val  
Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 39

atggccaagtacctggagctcgaagagggcggggtcataatgcaggcgttctactgggacgtgccttcaggaggaatatgggtggacacaat  
acggcagaagataccggagtggtagcatgccggaatctccgcaatatggattcctcccgagcaggggtatgagcggcggtattcgtatgg  
gctacgacctacgattattttgacctcggtgagtactaccagaagggaacgggtggaacgaggttcggctcaagcaggagctcataaacat  
gataaacaccgcccacgcctatggcatgaaggtaatagccgatatagtcatcaaccaccgcgccggcggtgacctggagtgaacccctcgt  
gaacgactatacctggaccgacttctcaaaggtcgcgtcgggtaatacacggccaactacctgacttccaccgaacgagctccatgcggg  
cgattccggaacatttggaggtatcccgacatatgccacgacaagagctgggaccagtactggctctggggccagccaggagctacgcgg  
catatctcaggagcatcgggtatgccttggcgctttgactacgtgaagggtacggagcgtgggtcgtcaaggactggctcaactgggtggg  
gcgggtggccgttggcgagtactgggaccccaacgttgatgccctcctccctgggcctactcgagcggcgccaaggctctcgaactccgcg  
tctactacaagtggatgaggcctttgacaacaaaacattccagcgctcgtctctgccttcagaacggccagactgtgtctcccgcgacccg  
ttcaaggccgtaacctttgtagccaaccacgataccgatataatctggaacaagtatccagcctacgcgttcacctcacctacgagggccagcc  
gacaatattctaccgcgactacgaggagtgggtcaacaaggataagctcaagaacctcatctggatacatgacaacctcggcgagggaagca  
ccgacatagtctactacgataaacgatgaactcatcttcgtaggaacgggtacggggacaagccggggcttataacctacatcaacctaggtc  
gagcaaggccggaagggtgggtctacgttcgaagttcgcgggagcgtgcatccacgagtagcaccggcaacctcggcggtgggtggacaa  
gtgggtggactcaagcgggtgggtgtacctgaggccctgccacgacccggccaacggctattacggctactccgtctggagctactgcg  
gggtgggctga

SEQ ID NO.: 40

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Arg Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala

Figure 16 (cont.)

Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Pro Asn Val Asp Ala Leu  
Leu Pro Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala  
Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg Asp  
Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr  
Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr  
Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu Tyr  
Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro  
Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 41

atggccaagtacctggagctcgaagagggcgggggtcataatgcaggcgttctactgggacgtgccttcaggaggaatatgggtgggacacaat  
acggcagaagataccggagtggtagatgccggaatctccgcaatatggattcctcccgagcaagggtatgagcggcggttattcgatgg  
gctacgacccctacgattattttgacctcggtgagtactaccagaagggaacgggtggaacgaggttcggctcaagcaggagctcataaacat  
gataaacacggccccatgcctacggcataaaggtcatagcggacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccgttc  
gttggggactacacctggacggacttctcaaaggtggcctcgggcaaatatactgccaactacctcactccacccgaacgagctccatgcg  
ggcgattccggaacatttggaggctatcccagacatagccacgacaagagctgggaccagtactggctctggggcaccaggagagctacgc  
ggcatatctcaggagcatcggcatcgcctggcgctttgactacgtgaagggctacggagcgtgggtcgtcaaggactggctcaactggtg  
ggcgcggtggcggttggcgagtactgggacaccaacgttgatgactcctcaactgggctactcagcggcgccaaggtcttcgactccc  
gcttactacaagatggacgcggcctttgacaacaagaacattcccgactcgtcgaggccctcaagaacgggggcacagtcgtcagccgcg  
acccgtttaagggcgtaaccttcgttgaaccacgacaccgatataatctggaacaagtatccagcctacgcgttcacctcacgagggc  
cagccgacaatatctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccggagg  
aagcacgagcatagtctactacgacagcgacgagatgatcttcgtgaggaacggctatggaagcaagcctggccttataacttacatcaacctc  
ggctcagcaaggttgaaggtgggtttatgtccgaagttcgcgggcgctgcatccacgagtatactggtaacctcgagggtgggtagac  
aagtacgtctactcaagcggctgggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgtgtggagctactgcg  
gtgttggtga

SEQ ID NO.: 42

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Ala  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Ser Ile Val Tyr  
Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

Figure 16 (cont.)

SEQ ID NO.: 43

atggccaagtactccgagctggaagagggcggttataatgcaggccttactgggacgtcccaggtggaggaatctggtgggacacat  
caggagcaagataccggagtggtagagggcggaataatccgccatttgattccccggcgagcaagggcatggcgggcgctattcgatg  
ggctacgacctacgacttcttgacctcggtagtagacgaccagaagggaaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaggctcatagcggacatcgtcataaaccaccgcgcaggcgagacctcagtggaacccg  
ttcgttggggactacacctggacggacttctcaaagggtggcctcgggcaataactgccaactacctcacttccaccccaacgaggtcaagt  
gctgtgacgagggcacatttgagggttcccagacatagcccacgagaagagctgggaccagcactggctctgggcgagcgatgagagcta  
cgccgctacctaaggagcatcggcgttgatgcctggcgcttcgactacgtcaagggtacggagcgtgggtcgtcaaggactggctggact  
gggtggggaggctggcgctggggagtagtgggacacaaacgttgatgactgctcaactgggctactcagcgatgcaaaagtcttcgac  
tccccgtctactacaagatggatgaggcctttgacaacaaaaacattccagcgctcgtctcgtccctcagaacggccagactgttctccgc  
gacccgttcaaggcgtaacctttgtagcaaacaccagacaccgatataatctggaacaagtatccagcctacgcgttcacctcacctacgagg  
gccagccgacaatattctaccgcgactacgaggagtggctcaacaaggataaagctcaagaacctcatctggatacatgacaacctcgtcggag  
gaagcagagcatagttactacgacagcgacgagatgatctcgtgaggaacggctatggaagcaagcctggccttataactacatcaacct  
cggctcgagcaaggttgaaggtgggtttacgttccgaagttcgaggtcgtgcatacacgagtagaccggcaatctcggcggtgggtgg  
acaagtgggtgactcaagcggctgggttacctcagggtcctgcccagacccggccaacggccagtagcggctactccgtctggagctac  
tgcggtgttgctga

SEQ ID NO.: 44

Met Ala Lys Tyr Ser Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp  
Val Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp  
Glu Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Val Gly Gly Ser Thr Ser Ile Val  
Tyr Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 45

atggccaagtactccgacctggaagagggcggttataatgcaggccttactgggacgtcccaggtggaggaatctggtgggacacat  
caggagcaagataccggagtggtagagggcggaataatccgccatttgattccccggcgagcaagggcatggcgggcgctattcgatg  
ggctacgacctacgacttcttgacctcggtagtagacgaccagaagggaaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaggctcatagcggacatcgtcataaaccaccgcgcaggcgagacctcagtggaacccg  
ttcgttggggactacacctggacggacttctcaaagggtggcctcgggcaataactgccaactacctcacttccaccccaacgaggtcaagt  
gctgtgacgagggcacatttgagggttcccagacatagcccacgagaagagctgggaccagcactggctctgggcgagcgatgagagcta  
cgccgctacctaaggagcatcggcgttgatgcctggcgctttgactacgtgaagggtacggagcgtgggtcgtcaaggactggctcaactg  
gtggggcggtggggcgttggcgagtactgggacaccaacgttgatgactcctcaactgggctactcagcggcgccaaggtcttcgactt  
cccgtctactacaagatggatgaggcctttgacaacaaaaacattccagcgctcgtctcgtccctcagaacggccagactgttctccgcg  
acccgttcaaggccgtaacctttgtagcaaacaccagacaccgatataatctggaacaagtatccagcctacgcgttcacctcacctacgagg  
ccagccgacaatattctaccgcgactacgaggagtggctcaacaaggataaagctcaagaacctcatctggatacatgacaacctcggcgag

Figure 16 (cont.)

gaagcaccgacatagtctactacgataacgatgaactcatcttcgtcaggaacggctacggggacaagccggggcttataacctacatcaacct  
aggctcgagcaaggccggaaggtgggtttatgtgccgaagttcgccggcgctgcacccaggtatactggttaacctcggaggctgggtg  
acaagtacgtctactcaagcggctgggtctatctcgaagctccagcttacgacctgccaacgggcagtagtgctactccgtgtggagctattgc  
gggtgtgggtga

SEQ ID NO.: 46

Met Ala Lys Tyr Ser Asp Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp  
Val Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp  
Glu Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val  
Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 47

atggccaagtacaccgagctggaagagggcgccgttataatgcaggccttctactgggacgtcccagggtggaggaatctggtgggacaccat  
caggagcaagataccggagtgtacgagggcggaatatccgccatttgattccccggcgagcaaggggcatggcgcgccatttcgatg  
ggctacgaccctacgacttctttgacctcgggtgagtacgaccagaaggggaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacaccgcccacgcctatggcatgaaggtaatagccgatatagtcatcaaccaccgcgccggcggtgacctggagtggaaacctt  
cgtgaacgactatacttgaccgacttctcaaaggctcgcgtcgggtaatacacggccaactacctcacttccacccaacgaggtcaagt  
ctgtgacgagggcacatttgaggcttcccagacatagcccacgagaagagctgggaccagcactggctctgggcgagcgatgagagctac  
gccgcctacctaaaggagcatcgcggttgatgcctggcgctttgactacgtgaagggtacggagcgtgggtcgtcaaggactggctcaactgg  
tggggcggttggggcgttggcgagtactgggacaccaacgttgatgcactcctcaactgggcctactcagcggcgccaaggtcttcgacttc  
ccgctctactacaagatggatgaggcctttgacaacaaaacattccagcgctcgtctctgccttcagaacggccagactgttgcctccgcga  
cccgttaaggccgtaaccttttagcaaacacacgacaccgatataatctggaacaagtaccttgcttatgctttcatctcactacgaaggcca  
gcccgtcatattctaccgcgactacgaggagtggctcaacaaggacaggtgaacaacctcatatggatacacgaccacctcgcaggtggaag  
cacgagcatagtattactacgacagcgacgagatgatcttcgtgaggaacggctatggaagcaagcctggccttataacttacatcaacctcggct  
cgagcaaggttgaaggtgggtttacgtccgaagttcgaggcccggtgcatacacgagtagaccggcaatctcggcggtgggtggacaag  
tgggtggactcaagcggctgggtctacctcgaggctcctgccacgaccggccaacggccagtaggctactccgtctggagctactcggg  
tgttgggtgag

SEQ ID NO.: 48

Met Ala Lys Tyr Thr Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys

Figure 16 (cont.)

Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp  
Val Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp  
Glu Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Leu  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Arg Leu Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Ser Ile Val  
Tyr Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Pro Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 53

ATGGCCAAGTACTCCGAGCTGGAAGAGGGCGGCGTTATAATGCAGGCCTTCTACTGGG  
ACGTCCCAGGTGGAGGAATCTGGTGGGACACCATCAGGAGCAAGATACCGGAGTGGT  
ACGAGGCGGGAATATCCGCCATTGGATTCCCCCGGCGAGCAAGGGCATGGGCGGCG  
CCTATTCGATGGGCTACGACCCCTACGACTTCTTTGACCTCGGTGAGTACGACCAGAA  
GGGAACGGTAGAGACGCGCTTTGGCTCCAAGCAGGAGCTCGTGAACATGATAAACAC  
GGCCCATGCCTACGGCATAAAGGTCATAGCGGACATCGTCATAAACCACCGCACAGGC  
GGAGACCTCGAGTGGAACCCGTTCTGGTGGGACTACACCTGGACGGACTTCTCAAAGG  
TGGCCTCGGGCAAATATACTGCCAACTACCTCGACTTCCACCCCAACGAGGTCAAGTG  
CTGTGACGAGGGCACATTTGGAGGCTTCCCAGACATAGCCACGAGAAGAGCTGGGA  
CCAGCACTGGCTCTGGGCGAGCGATGAGAGCTACGCCGCTACCTAAGGAGCATCGGC  
GTTGATGCCTGGCGCTTCGACTACGTCAAGGGCTACGGAGCGTGGGTCGTCAAGGACT  
GGCTGGACTGGTGGGGAGGCTGGGCCGTCGGGGAGTACTGGGACACAAACGTTGATG  
CACTGCTCAACTGGGCCTACTCGAGCGATGCAAAAAGTCTTCGACTTCCCGCTCTACTAC  
AAGATGGATGAGGCCTTTGACAACAAAAACATTCCAGCGCTCGTCTCTGCCCTTCAGA  
ACGGCCAGACTGTTGTCTCCCGCGACCCGTTCAAGGCCGTAACCTTTGTAGCAAACCA  
CGACACCGATATAATCTGGAACAAAGTATCCAGCCTACGCGTTCATCCTCACCTACGAG  
GGCCAGCCGACAATATTCTACCGCGACTACGAGGAGTGGCTCAACAAGGATAAGCTCA  
AGAACCTCATCTGGATACATGACAACCTCGCCGGAGGAAGCACTGACATCGTTTACTA  
CGACAACGACGAGCTGATATTCTGTGAGAAACGGCTACGGAAGCAAGCCGGGACTGAT  
AACATACATCAACCTCGCCTCAAGCAAAGCCGGAAGGTGGGTCTACGTTCCGAAGTTC  
GCGGGAGCGTGCATCCACGAGTACACCGGCAACCTCGGCGGCTGGGTGGACAAGTGG  
GTGGACTCAAGCGGGTGGGTGTACCTCGAGGCCCTGCCACGACCCGGCCAACGGCT  
ATTACGGCTACTCCGTCTGGAGCTACTGCGGTGTTGGCTGA

SEQ ID NO.: 54

MAKYSELEEGGVIMQAFYWDVPGGGIWWDTIRSKIPEWYEAGISAIWIPPASKGMGGAYS  
MGYDPYDFDLGEYDQKGTVETRFGSKQELVNMINAHAYGIKVIADIVINHRTGGDLEW  
NPFVGDYTWTDVFSKVASGKYTANYLDFHPNEVKCCDEGTFGGFPDIAHEKSWDQHWLW  
ASDESYAAYLRSIGVDAWRFDYVKGYGAWVVKDWLDWWGGWAVGEYWDTNVDALL  
NWAYSSDAKVDFDPLYKMDAEDFNKNIPALVSALQNGQTVVSRDPFKA VTFVANHDTD  
IWNKYPAYAFILTYEGQPTIFYRDYEEWLNKDKLKNLIWIHDNLAGGSTDIVYYDNDELIF  
VRNGYGSKPGLITYINLASSKAGRWWVYPKFAGACIHEYTGNLGGWVDKWVDSSGWVY  
LEAPAHDPANGYYGYSVWSYCGVG

SEQ ID NO.: 55

Figure 16 (cont.)

ATGGCCAAGTACCTGGAGCTCGAGGAGGGCGGGGTCATAATGCAGGCGTTCTACTGGG  
ACGTGCCTTCAGGAGGAATATGGTGGGACACAATACGGCAGAAGATACCGGAGTGGT  
ACGATGCCGGAATCTCCGCAATATGGATTCCCCCGGCGAGCAAGGGCATGGGCGGCGC  
CTATTCGATGGGCTACGACCCCTACGACTTCTTTGACCTCGGTGAGTACGACCAGAAG  
GGAACGGTAGAGACGCGCTTTGGCTCCAAGCAGGAGCTCGTGAACATGATAAACACC  
GCCCACGCCTATGGCATGAAGGTAATAGCCGATATAGTCATCAACCACCGCGCCGGCG  
GTGACCTGGAGTGGAACCCCTTCGTGAACGACTATACCTGGACCGACTTCTCAAAGGT  
CGCGTCGGGTAAATACACGGCCAACTACCTCGACTTCCACCCGAACGAGCTCCATGCG  
GGCGATTCCGGAACATTTGGAGGCTATCCCGACATATGCCACGACAAGAGCTGGGACC  
AGTACTGGCTCTGGGCCAGCCAGGAGAGCTACGCGGCATATCTCAGGAGCATCGGCAT  
CGATGCCTGGCGCTTTGACTACGTGAAGGGCTACGGAGCGTGGGTCTGTCAGGACTGG  
CTCAACTGGTGGGGCGGCTGGGCCGTTGGCGAGTACTGGGACACCAACGTTGATGCAC  
TCCTCAACTGGGCCTACTCGAGCGGCGCCAAGGTCTTCGACTTCCCGCTCTACTACAAG  
ATGGATGAGGCCTTTGACAACAAAAACATTCCAGCGCTCGTCTCTGCCCTTCAGAACG  
GCCAGACTGTTGTCTCCCGCGACCCGTTCAAGGCCGTAACCTTTGTAGCAAACCACGA  
CACCGATATAATCTGGAACAAGTACCTTGCTTATGCTTTCATCCTCACCTACGAAGGCC  
AGCCCGTCATATTCTACCGCGACTACGAGGAGTGGCTCAACAAGGACAGGTTGAACAA  
CCTCATATGGATACACGACCACCTCGCAGGTGGAAGCACGAGCATAGTTTACTACGAC  
AGCGACGAGATGATCTTCGTGAGGAACGGCTATGGAAGCAAGCCTGGCCTTATAACTT  
ACATCAACCTCGGCTCGAGCAAGGTTGGAAGGTGGGTTTACGTTCCGAAGTTCGCAGG  
CTCGTGCATACACGAGTACACCGGCAATCTCGGCGGCTGGGTGGACAAGTGGGTGGAC  
TCAAGCGGCTGGGTCTACCTCGAGGCTCCTGCCACGACCCGGCCAACGGCCAGTACG  
GCTACTCCGTCTGGAGCTATTGCGGTGTTGGCTGA

SEQ ID NO.: 56

MAKYLELEEGGVIMQAFYWDVPSGGIWWDITRQKIPEWYDAGISAIWIPPASKGMGGAYS  
MGYDPYDFFDLGEYDQKGTVETFRGSKQELVNMINTAHAYGMKVIADIVINHRAGGDLE  
WNPVNDYTWTDFSKVASGKYTANYLDFHPNELHAGDSGTFGGYPDICHDKSWDQYWL  
WASQESYAAYLRSIGIDAWRFDYVKGYGAWVVKDWLNWWGGWAVGEYWDTNVDALL  
NWAYSSGAKVDFPLYYKMDEAFDNKNIPALVSALQNGQTVVSRDPFKAVTFVANHDTD  
IWNKYLAYAFILTYEQPVIFYRDYEEWLNKDRLNNLIWIHDHLAGGSTSIVYYDSDEMIF  
VRNGYGSKPGLITYINLGSSKVGRWVYVPKFAGSCIHEYTGNLGGWVDKWVDSSGWVYL  
EAPAHDPANGQYGYSVWSYCGVG

SEQ ID NO.: 57

ATGGCCAAGTACCTGGAGCTCGAAGAGAGCGGGGTCATAATGCAGGCGTTCTACTGGG  
ACGTGCCTTCAGGAGGAATATGGTGGGACACAATACGGCAGAAGATACCGGAGTGGT  
ACGATGCCGGAATCTCCGCAATATGGATTCTCCCGCGAGCAAGGGTATGAGCGGCGG  
CTATTCGATGGGCTACGACCCCTACGATTATTTGACCTCGGTGAGTACTACCAGAAGG  
GAACGGTGGAACGAGGTTTCGGCTCAAAGCAGGAGCTCATAAACATGATAAACACCG  
CCCACGCCTACGGCATCAAGGTCATCGCAGACATAGTAATCAACCACCGCGCCGGAGG  
AGACCTTGAGTGGAACCCCTTCGTCAATGACTACACCTGGACGGACTTCTCGAAGGTC  
GCTTCCGGCAAGTACACGGCCAACTACCTCGACTTCCACCCCAACGAGGTCAAGTGCT  
GTGACGAGGGGCACATTTGGAGGCTTCCAGACATAGCCACGAGAAGAGCTGGGACC  
AGCACTGGCTCTGGGCGAGCGATGAGAGCTACGCCGCCTACCTAAGGAGCATCGGCGT  
TGATGCCTGGCGCTTTGACTACGTGAAGGGCTACGGAGCGTGGGTCTGTCAGGACTGG  
CTCAACTGGTGGGGTGGCTGGGCCGTCGGGGAGTACTGGGACACAAACGTTGATGCAC  
TGCTCAACTGGGCCTACTCGAGCGATGCAAAAAGTCTTCGACTTCCCGCTCTACTACAAG  
ATGGACGAGGCCTTCGATAACAACAACATTCCCGCCCTGGTGGACGCCCTCAGATACG  
GTCAGACAGTGGTCAGCCGCGACCCGTTCAAGGCTGTGACGTTTGTAGCCAACCACGA

Figure 16 (cont.)

TACCGATATAATCTGGAACAAGTACCTTGCTTATGCTTTCATCCTCACCTACGAAGGCC  
AGCCCGTCATATTCTACCGCGACTACGAGGAGTGGCTCAACAAGGACAGGTTGAACAA  
CCTCATATGGATACACGACCACCTCGCAGGTGGAAGCACTGACATCGTTTACTACGAC  
AACGACGAGCTGATATTCGTGAGAAACGGCTACGGAAGCAAGCCGGGACTGATAACA  
TACATCAACCTCGCCTCAAGCAAAGCCGGAAGGTGGGTCTACGTTCCGAAGTTCGCGG  
GAGCGTGCATCCACGAGTACACCGGCAACCTCGGCGGCTGGGTGGACAAGTGGGTGG  
ACTCAAGCGGGTGGGTGTACCTCGAGGCCCTGCCACGACCCGGCCAACGGCTATTA  
CGGCTACTCCGTCTGGAGCTATTGCGGTGTTGGCTGA

SEQ ID NO.: 58

MAKYLELEESGVIMQAFYWDVPSGGIWWDTIRQKIPEWYDAGISAIWIPPASKGMSGGYS  
MGYDPYDYFDLGEYYQKGTVETFRGSKQELINMINTAHAYGIKVIADIVINHRAGGDLEW  
NPFVNDYTWTDFSKVASGKYTANYLDFHPNEVKCCDEGTFGGFPDIAHEKSWDQHWLW  
ASDESYAAYLRSIGVDAWRFDYVKGYGAWVVKDNLNWWGGWAVGEYWDTNVDALL  
NWAYSSDAKVDFDPLYKMDFAFDNNIPALVDALRYGQTVVSRDPFKA VTFVANHDTD  
IWNKYLAYAFILTYEGQPVIFYRDYEEWLNKDRLNLIWIHDHLAGGSTDIVYYDNDELIF  
VRNGYGSKPGLITYINLASSKAGRWWYVPKFAGACIHEYTGNLGGWVDKWVDSSGWVY  
LEAPAHDPANGYYGYSVWSYCGVG

SEQ ID NO.: 59

atggccaagtacgtgagctcgaagagggcggggtcataatgcaggcgttctactgggacgtgccttcaggaggaatatgggtggacacaat  
acggcagaagataccggagtgtacgatgccgaatctccgaatatgattcctcccgagcaagggatgagcggcggtattcgtatgg  
gtacgacccctacgattatttgacctcggtgagtactaccagaaggggaacgggtggaacgaggttcggctcaaagcaggagctcataacat  
gataaacaccgcccacgctacggcatcaaggtcatcgcagacatagtaataaccaccgcccggaggagaccttgagtgaacccttcg  
tcaatgactacacgtgacggacttctgaaggtcgttcggcaagtacacggccaactacctcgactccaccgaacgagctccatcggg  
gcgattccggaacatttggaggctatcccgacatatgccacgacaagagctgggaccagtactggctctgggccagccaggagagctacgcg  
gcatactcaggagcatcgccatcgatgcctggcgctcgactacgtcaagggctatgctccctgggtcgtaaggactggctgaactgggtggg  
gaggctgggcgggttgagagtagtgggacaccaacgtcgacgctgttctcaactgggcatactcgagcgggtccaaggtcttgacttcgcct  
ctactacaagatggacgagccttcgataacaacaacattcccgccctggtagccctcagatacggtcagacagtggtcagccgcgacc  
cgttcaaggctgtgacgttttagccaaccacgataccgatataatttgaacaagtacccggcctacgccttcacctacgagggccag  
ccgacgatatctaccgcgactacgaggagtgggtcaacaaggacagggtcaagaacctcatctggatacacgaccacctgcgggtggaag  
cactgacatcgttactacgacaacgacgagctgatattcgtgaaacgggtacggaaagcaggcggactgataacatacatcaacctgc  
gtcaagcaaagccggaaggtgggttatgtgccgaagttcgcgggcgctgcatccacgagtatactggtaacctcgagggtgggtgagaca  
agtactgtactcaagcggctgggtctatctgaagctccagcttacaccctgccaacgggcagtaggtactccgtgtggagctattcggt  
gttggtga

SEQ ID NO.: 60

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val  
Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu Ala  
Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp  
Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr

Figure 16 (cont.)

Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Arg Leu Lys Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr  
Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Ala Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu Tyr  
Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro  
Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 61

atggccaagtactccgagctgaaaaagggcggggtcataatgcaggcgttctactgggacgtgccttcaggaggaatatggtgggacacaat  
acggcagaagataccggagtggtagagggcggaatatccgccatttggattctcccgcgagcaagggtatgagcggcggtattcgcgatg  
gctacgacccctacgattatttggacctcggtagtactaccagaaggggaacgggtgaaacgaggttcggctcaaagcaggagctcataacat  
gataaacaccgcccacgcctacggcatcaaggtcatcgcagacatagtaataaccaccgcgcggaggagaccttgagtggaaaccccttcg  
tcaatgactacacctggacggacttctcgaaggtcgttccggcaagtacacggccaactaccctaactccaccgaaacgagctccatgcgg  
gcgattccggaacatttggagggtatcccgcacatagccacgacaagagctgggaccagtactggctctgggccagccaggagagctacgcg  
gcatactcaggagcatcggcatcgatgcctggcgcttcgactacgtcaagggctacggagcgtgggtcgtcaaggactggctggactgggtg  
gggaggtcgggcccgtcggggagtactgggacacaaacgttgatgcactgctcaactgggcctactcgagcgtgcaaaagcttcgacttccc  
gctctactacaagatggatgaggcctttgacaacaaaaacattccagcgctcgtctctgccttcagaacggccagactgttctctcccgcgacc  
cgttaaggccgtaaccttttagcaaacatgacaccgatataatctggaacaagtatccagcctacgcgttcactcctacacacgagggccag  
ccgacaatattctaccgcgactacgaggagtgggtcaacaaggataagctcaagaacctcatctggatacatgacaacctcggcggagggaag  
caccgacatagctactacgataacgatgaactcatctcgtcaggaacgggtacggggacaagccggggttataacctacatcaacctaggc  
tcgagcaaggccggaaggtgggtctacgtccgaagttcgcgggagcgtgcacccaggtacacccggcaacctcggcgggtgggtggaca  
agtgggtggactcaagcgggtgggtgtacctcagggccctgcccacgacccggccaacggctattacggctactccgtctggagctactgc  
ggggtgggctga

SEQ ID NO.: 62

Met Ala Lys Tyr Ser Glu Leu Lys Lys Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asn Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val  
Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu  
Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr  
Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 63

atggccaagtactccgagctgaaagagggcggggtcataatgcaggcgttctactgggacgtgccttcaggaggaatatggtgggacacaat  
acggcagaagataccggagtggtagatgccggaatctccgcaatatggattccccggcgagcaagggtatggcggcgcttattcgcgatg  
ggctacgacccctacgacttctttgacctcggtagtactaccagaaggggaacggtagagacgcgctttggctccaagcaggagctcgtgaa  
catgataaacacggcccatgcctacggcataaaggccatagcggacatcgtcataaaccaccgcgcaggcggagacctcagtggaacccg

Figure 16 (cont.)

ttcgttggggactacacctggacggacttctcaaaaggtggcctcgggcaaatactgccaactacctcgacttccacccaacgaggtcaagt  
gctgtgacgagggcacatttggaggcttcccagacatagccacgagaagagctgggaccagcactggctctgggcgagcgatgagagcta  
cgccgcctacctaagggagcattggcggtgatgcctggcgcttgactacgtgaagggtacggagcgtgggtcgtaaggactggctcaactg  
gtggggcggtggggcggtggcgagtgactgggacaccaacgttgatgcactcctcaactgggcctactcgagcggcgccaaggtcttcgactt  
cccgtctactacaagatggacgcggcctttgacaacaagaacattcccgcactcgtcgaggccctcaagaacgggggcacagtcgtcagcc  
gcgacccgtttaaggccgtaaccttcgttgcaaacacgacaccgatataatctggaacaagtatccagcctacgcgttcacctacacacgag  
ggccagccgacaatattctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccgg  
aggaagcaccgacatagctactacgataacgatgaactcatcttcgtcaggaacggctacggggacaagccggggcttataacctacatcaa  
cctaggctggagcaaggccggaaggtgggtttatgtgccgaagttcggggcggtgcacccagagtatactggtaacctcggaggctggg  
tagacaagtacgtctactcaagcgggtggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgtgtggagcta  
ctgcgggggtgggggtga

SEQ ID NO.: 64

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp  
Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val Asn  
Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Ala Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly  
Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys  
Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly  
Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr  
Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp  
Val Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp  
Ala Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp  
Ala Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val Ser  
Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro  
Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu  
Asn Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val  
Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr  
Ile Asn Leu Gly Trp Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His  
Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu Glu  
Ala Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 65

atggccaagtactccgagctggaagaaggcggcggttataatgcaggccttctactgggacgtcccagggtggaggaatctggtggggcaccat  
caggagcaagataccggagtggtacgagggcggaatatccgccatttggattcctcccgcgagcaagggtatgagcggcggtattcgtatgg  
gctacgacccctacgattattttgacctcgggtgagtactaccagaagggaacgggtgaaacgaggttcggctcaaagcaggagctcataaacat  
gataaacaccgcccacgcctatggcatgaaggtaatacgcatatagtcataaccaccgcggcggtgacctggagtgaaccccttcgt  
gaacgactatacctggaccgacttctcaaaggctcgcgtcggttaatacacggccaactacctcgacttccacccgaacgagctccatgcggg  
cgattccggaacatttggaggctatcccacatatgccacgacaagagctgggaccagtgactggctctggccagccaggagagctacgagg  
catatctcaggagcatcggcatcgtatgcttgcgttcgactacgtcaagggtatgctccctgggtcgtcaaggactggctgaactggtgggg  
aggctgggcgggttgagagtactgggacaccaacgtcgacgtgttcacttgggcatactcgagcgggtgccaaaggtctttgacttcgccctc  
tactacaagatggacgagggccttcgataacaacaacattcccgccttgggtggacgccctcagatacggtcagacagtggtcagccgcgacc  
gttcaaggctgtgacgtttgtagccaaccacgataccgataaatttgaacaagtacccggcctacgccttcacctacacgaggccagc  
cgacgatattctaccgcgactacgaggagtggctcaacaaggacaggctcaagaacctcatctggatacacgaccacctcgccggttgaagc  
acgagcatagttactacgacagcgacgagatgatcttcgtgaggaaacggctatggaagcaagcctggccttataacttacatcaacctcggctc  
gagcaagggttgaaggtgggttacgtccgaagttcgcaggctcgtgcatacacgagtagacacggcaatctcggcggttgggtggacaagt  
gggtggactcaagcggttgggtctacctcagggtcctgcccacgacccggccaacggccagtagggctactccgtctggagctatttcggt  
gttggctga

Figure 16 (cont.)

SEQ ID NO.: 66

Met Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly  
Gly Gly Ile Trp Trp Gly Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala Ile Trp Ile  
Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp Leu  
Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met Ile  
Asn Thr Ala His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr  
Pro Asp Ile Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val  
Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val  
Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu Ala  
Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp  
Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr  
Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Arg Leu Lys Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Ser Ile Val Tyr Tyr  
Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr  
Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro  
Ala His Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 67

atggccaagtacctggagctcgaagaggcggtgcataatgcaggcgttctactgggacgtgccttcgggaggaatatggtgggacacaat  
acggcagaagataccggagtgtacgatcccggaatctccgcaatatgattcctcccgcgagcaagggtatgagcggcggtattcgatgg  
gctacgaccctacgattattttgacctcggtagtactaccagaagggaacgggtgaaacgaggttcggctcaaagcaggagctcataaacat  
gataaacacggcccatgcctacggcataaaggtcatagcggacatcgtcataaacaccgcgcaggcggagacctcagtggaacccgttc  
gttggggactacacctggacggacttctcaaaggtggcctcgggcaaatatactgccaaactacctcgaactccacccaacagagtgcaagtgtc  
gtgacgagggcacatttgagggttcccagacatagcccacgagaagagctgggaccagcactggctctgggcgagcagatgagagctacg  
ccgctacctaaggagcatcggcggtgatgcctggcgcttcgactacgtcaagggtcagcagcgtgggtcgtcaaggactggctggactgtt  
ggggaggtcgggcccgtcggggagtactgggacacaaacgttgatgcactgtcaactgggctactcgcagcagatgcaaaagtcttcgacttc  
ccgcttactacaagatggacgagggccttcgataacaacaacattcccgccttggtggacgccctcagatacggctcagacagtgtcagccgc  
gaccggttcaaggctgtgacgtttgtagccaaccacgataccgatataatctggaacaagtatccagcctacgcgttcactcactacgaggg  
ccagccgacaatatctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctcgccggag  
gaagcagcagcatagttactacgacagcagcagatgatcttcgtgaggaacggctatggaagcaagcctggcctataacttacatcaacct  
cggctcgagcaagggtggaaggtgggtctacgttcgaagttcgcgggagcgtgcatccacgagtagaccggcaacctcgccgggtggtg  
gacaagtgggtggactcaagcgggtgggtgtacctcgaggccctgcccacgaccggccaacggctattacggctactccgtctggagcta  
ctgcgtggtgggctga

SEQ ID NO.: 68

Met Ala Lys Tyr Leu Glu Leu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Ser  
Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp  
Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp  
Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met  
Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe Gly Gly  
Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser Tyr Ala  
Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val  
Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala  
Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu

Figure 16 (cont.)

Ala Phe Asp Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg  
Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala  
Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn  
Lys Asp Lys Leu Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Ser Ile Val Tyr  
Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile  
Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala  
Pro Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Val Val Gly

SEQ ID NO.: 73

atggctctggaagaggggcggttataatgcaggccttactgggacgtcccagggtggaggaatctggtgggacacatagcccagaagat  
accgactgggcgagcgccgggatttcggcaatatggattcctcccgcgagtaagggcagcggcggtattcgatgggctacgaccct  
acgattcttcgacctcggtgagtactaccagaagggaagcgttgagaccgcttcggatcaaaaggagcgttgtaacatgataaacaccgc  
ccatgctcacaacatgaaggtcatagcggacatagtcacaccacgcgcggcgacgtggagtgggaatccttaccacacagctacac  
ctggaccgatttctgaaggtcgcgtcgggcaagtacacggccaactacctgacttccacccgaacgagcttcacgcggcggtattccggaa  
catttggaggctatcccgacatagccacgacaagagctgggaccagcactggctctgggccaagcaagaaagctacgcgcctacccggg  
agcatcggcatcgacgcctggcgcttcgactacgtcaagggctacgctcctgggtcgttaagaactggctgaaccgggtggggcggtgggc  
ggttggagagtactgggacaccaacgtcgtacgactcctgagctgggctacgacagcggtgctaaagtcttcgacttcccgtctactacaag  
atggacgaggccttcgataacaacaacatccccgccctcgtggacgccctcaagaacggaggcacggctcagccgcgacccgttcaag  
ccgtgaccttcgttgcaaccacgataccaacataatctggaacaagatccggcctacgccttcacctacacatgagggacagccggcaat  
attctaccgcgactacgaggagtggctcaacaaggacaggtcaggaacctcatctggatacacgaccacctcgcgggagggaagcacagac  
atcatctactacgacagcgacgagcttatctcgtgaaacggctacggggacaagccgggactgataacctacatcaacctcggctcaagc  
aaggccggaaggtgggtctactgctcgaagttcgcaggctcgtgcatacacgagtagaccggcaacctcggcggtggattgacaagtggtg  
tgactcaagcggctcgggtctaccttgaggccccgccacgacccggccaacggccagtagggctactccgtatggagctactcgggtgttg  
ggtga

SEQ ID NO.: 74

Met Ala Leu Glu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly Gly Gly Ile Trp  
Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala  
Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp Leu Gly Glu Tyr  
Tyr Gln Lys Gly Ser Val Glu Thr Arg Phe Gly Ser Lys Glu Glu Leu Val Asn Met Ile Asn Thr Ala  
His Ala His Asn Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp Leu Glu Trp  
Asn Pro Phe Thr Asn Ser Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn  
Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr Pro Asp Ile  
Cys His Asp Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asn Glu Ser Tyr Ala Ala Tyr Leu Arg  
Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val Lys Asn Trp  
Leu Asn Arg Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Leu Leu Ser Trp  
Ala Tyr Asp Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp  
Asn Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Lys Asn Gly Gly Thr Val Val Ser Arg Asp Pro Phe  
Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asn Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile  
Leu Thr Tyr Glu Gly Gln Pro Ala Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Arg Leu  
Arg Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Ile Tyr Tyr Asp Ser Asp Glu  
Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Gly Ser Ser  
Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr Thr Gly Asn Leu  
Gly Gly Trp Ile Asp Lys Trp Val Asp Ser Ser Gly Arg Val Tyr Leu Glu Ala Pro Ala His Asp Pro  
Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 75

atggctctggaagaggggcggttataatgcaggcatttactgggacgtcccatgggaggaatctggtgggacacgatagcccagaagat  
accgactgggcaagcgccgggatttcggcgatatggattccccccgcgagcaagggtatgagcggcggtattcgatgggctacgaccct

Figure 16 (cont.)

acgattattttgacctcggtagtactaccagaaggggaacgggtggaacaagattcgggtcaaagcaggagctcataaacatgataaacaccg  
cccacgcctatggcatgaaggtgaatagccgatatagtcatcaaccaccgcgcggcggtgacttggagtggaaaccccttcgtgaacgactata  
cctggaccgacttctgaaggtcgcgtcgggtaaatacacggccaactacctcgacttccaccgaacgagctccacgcgggagattccgga  
acatttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggccagccaggagagctacgcggcctatctcag  
gagcatcggcatcgacgcctggcgttcgactacgtcaagggctatgctccctgggtcgtcagggactggctgaactgggtgggagggtggg  
cagttggagagtactgggacaccaacgtcgacgtgttctcaactgggcatactcgagcgggtccaaggtctttgacttcgccctctactacaag  
atggacgaggccttcgataacaacaacattcccgcctggtggacgccctcagatacggccagacagtgggtcagccgcgacccgttcaaggc  
tgtgacgttttagccaaccacgataccgacataatctggaacaagtatccagcctacgcgttcacctacgagggccagccgacaatat  
tctaccgcgactacgaggagtgggtcaacaaggacaagctcaagaacctcatctggatacatgacaacctgcgggaggaggagcactgacatc  
gtttactacgacaacgacgagctgatattcgtgagaaacggctacggaagcaagccgggactgataacatacatcaacctcggtcaagcaaa  
gccggaaggtgggtttacgttcgaagttcgcaggctcgtgcatacacgagtagaccggcaacctcgcggtgggtgggacaagtggtggga  
ctcaagcggctgggtttacctcgaggctcctgccacgacccggccaacggccagtagcggtactccgttggagctattgcggtgttgggtga

SEQ ID NO.: 76

Met Ala Leu Glu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met Gly Gly Ile Trp  
Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala  
Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp Leu Gly Glu Tyr  
Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met Ile Asn Thr Ala  
His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp Leu Glu Trp  
Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn  
Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr Pro Asp Ile  
Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala Tyr Leu Arg  
Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val Arg Asp Trp  
Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val Leu Asn Trp  
Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp Asn  
Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp Pro Phe Lys  
Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile  
Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Lys Leu  
Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr Asp Asn Asp  
Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Gly Ser Ser  
Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr Thr Gly Asn Leu  
Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro Ala His Asp Pro  
Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 77

atggctctggaagagggcggtcataatgcaggccttactgggacgtcccatgggaggaatctgggtgggacacgatagcccagaagat  
acccgactgggcaagcgccgggatttggcgatatggatccctcccgagcaagggtatgagcggcggtattcgatgggctacgaccct  
acgattattttgacctcggtagtactaccagaaggggaacgggtggaacagaggttcgggtcaaagcaggagctcataaacatgataaacaccg  
cccacgcctatggcatgaaggtgaatagccgatatagtcatcaaccaccgcgcggcggtgacctggagtggaaaccccttcgtgaacgactata  
cctggaccgacttctcaaggtcgcgtcgggtaaatacacggccaactacctcgacttccaccgaacgagctccatcggggagattccggaa  
catttggaggctatcccacatatgccacgacaagagctgggaccagtactggctctgggccagccaggagagctacgcggcctatctcagg  
agcatcggcatcgatgcctggcgttcgactacgtcaagggctatgctccctgggtcgtcaaggactggctgaactgggtgggagggtgggc  
gggtggagagtactgggacaccaacgtcgacgtgttctcaactgggcatactcgagcgggtccaaggtctttgacttcgccctctactacaaga  
tggacgaggccttcgataacaacaacattcccgcctggtggacgccctcagatacgggtcagacagtgggtcagccgcgacccgttcaaggct  
gtgacgttttagccaaccacgataccgacataatctggaacaagtatccagcctacgcgttcacctacgagggccagccgacaatat  
ctaccgcgactacgaggagtggctcaacaaggataagctcaagaacctcatctggatacatgacaacctgcgggaggaggagcactgacatcg  
tttactacgacaacgacgagctgatattcgtgagaaacggctacggaagcaagccgggactgataacatacatcaacctgcctcaagcaag  
ccggaaggtgggtttacgttcgaagttcgcaggctcgtgcatacacgagtagaccggcaatctcgcggtctgggtgggacaagtggtgggac  
tcaagcggctgggtctacctcgaggctcctgccacgacccggccaacggccagtagcggtactccgtctggagctactgcggtgttgggtga

Figure 16 (cont.)

SEQ ID NO.: 78

Met Ala Leu Glu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Met Gly Gly Ile Trp  
Trp Asp Thr Ile Ala Gln Lys Ile Pro Asp Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala  
Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp Leu Gly Glu Tyr  
Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile Asn Met Ile Asn Thr Ala  
His Ala Tyr Gly Met Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp Leu Glu Trp  
Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn  
Tyr Leu Asp Phe His Pro Asn Glu Leu His Ala Gly Asp Ser Gly Thr Phe Gly Gly Tyr Pro Asp Ile  
Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Gln Glu Ser Tyr Ala Ala Tyr Leu Arg  
Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Ala Pro Trp Val Val Lys Asp Trp  
Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Val Leu Asn Trp  
Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Ala Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp Asn  
Asn Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp Pro Phe Lys  
Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile  
Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Lys Leu  
Lys Asn Leu Ile Trp Ile His Asp Asn Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr Asp Asn Asp  
Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Ala Ser Ser  
Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr Thr Gly Asn Leu  
Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro Ala His Asp Pro  
Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 79

atgaagcctgcgaaactcctcgtctttgtgctcgtagtctctatcctcgcggggctctacgccagcccgcggggcgcccaagtacctggagc  
tcgaagaggggcggcgcataatgcagggcgttctactgggacgtgccttcaggaggaatatggtgggacacaatacggcagaagataccgga  
gtggtacgatgccggaatctccgcaatatggattccccggcgagcaaggcgatgggcggcgctattcgatgggtacgaccctacgactt  
ctttgacctcggtgagtagcaccagaagggaacggtagagacgcgctttggctccaagcaggagctcgtgaacatgataaacaccgccacg  
cctacggcatcaaggatcatcgagacatagtaataaccaccgcgcggaggagacctgagtgaaccccttcgtaatgactacacctgga  
cggacttctgaaggctcgttcggcaagtagcacggccaactacctgacttccacccaacgaggtcaagtgtcgcacgagggcacctttg  
gaggggtcccgacatagcccacgagaagagctgggaccagtactggctctgggcgagcaacgagagctacgccgctacctcaggagca  
tcggcggttacgcatggcgcttcgactacgtcaagggtacggagcgtgggtcgtcaaggactggctggactgggtgggaggctgggctg  
cggggagtactgggacacaaacgttgatgcactgtcactgggcctactcgagcgtatgcaaaagtcttcgacttcccgctctactacaagatg  
gacgggcctttgacaacaagaacattcccgactcgtcaggccctcaagaacgggggcacagtcgtcagccgcgacccgtttaaggccgt  
aaccttcgttgcaaacacgacacggacataattggaacaagtaccggcgctacgccttcacctacgagggccagccgacgatattc  
taccgcgactacgaggagtggctcaacaaggacaggctcaagaacctcatctggatacacgaccacctcgccggtggaagcaccgacatag  
tctactacgataacgatgaactcatcttcaggaacggctacggggacaagccggggttataacctacatcaacctaggctcgagcaagg  
ccgggaggtgggtctacgttcgaagttcggggagcgtgcattccacgagtagaccggcaacctcggcggtgggtggacaagtgggtgga  
ctcaagcgggtgggtgtacctcaggcccttcccacgaccggccaacggctattacggctactccgtctggagctactgcggggtgggct  
ga

SEQ ID NO.: 80

Met Lys Pro Ala Lys Leu Leu Val Phe Val Leu Val Val Ser Ile Leu Ala Gly Leu Tyr Ala Gln Pro  
Ala Gly Ala Ala Lys Tyr Leu Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val  
Pro Ser Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala  
Ile Trp Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe  
Phe Asp Leu Gly Glu Tyr Asp Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Val  
Asn Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly  
Gly Asp Leu Glu Trp Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly  
Lys Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe  
Gly Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Asn Glu Ser  
Tyr Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala

Figure 16 (cont.)

Trp Val Val Lys Asp Trp Leu Asp Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val  
Asp Ala Leu Leu Asn Trp Ala Tyr Ser Ser Asp Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met  
Asp Ala Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Glu Ala Leu Lys Asn Gly Gly Thr Val Val  
Ser Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr  
Pro Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Thr Ile Phe Tyr Arg Asp Tyr Glu Glu Trp  
Leu Asn Lys Asp Arg Leu Lys Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile  
Val Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Asp Lys Pro Gly Leu Ile Thr  
Tyr Ile Asn Leu Gly Ser Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile  
His Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Trp Val Asp Ser Ser Gly Trp Val Tyr Leu  
Glu Ala Pro Ala His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 81

atgaagaagttgtcgccctgttcataacccatgttttcgtagtgcagcatggcagctggtgcacagccagctagcgccgcaaagtattccgagctc  
gaagaaggcggcggtataatgcaggcccttactgggagctccaggtggaggaatctggtgggacaccatcaggagcaagataccggagt  
ggtacgagggggaataatccgccatttgattccgccagccagcaaggggatgagcggcggtactcgtatgggtacgatccctacgatttct  
tgacctcggcgagtacaaccagaagggaacctcgaaacgcgtttggtctaaacaggagctcatatgataaacacggcccatgacct  
cggcataaaggatcatagcggacatgctcataaaccaccgcgcaggcggagacctcagtggaacccgttcgttggggactacacctggacg  
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gcttccagacatagcccacgagaagagctgggaccagcactggctctgggcgagcgtatgagagctacggcgctacctaaggagcatcg  
gcgttgatgcctggcgctttgactacgtgaagggtctacggagcgtgggtcgtcaaggactggctcaactgggtggggcggtgggctggc  
gagtactgggacaccaacgttgatgcactcctcaactgggcctactcgagcggcgccaaggtcttcgacttcccgtctactacaagatggatg  
aggcctttgacaacaaaaacattccagcgctcgtctcgtccctcagaacggccagactgtgtctcccgcgacccgttcaaggccgtaaccttt  
gtagcaaacacgacaccgatataatctggaacaagtaccttgcttatgcttcatcctcacctacgaaggccagcccgtcatattctaccgcgac  
tacgaggagtggctcaacaaggacaggttgaaacaacctcatatggatacagaccacctcgaggtggaagcagcagcatagtctactacga  
cagcgacgagatgatcttcgtgaggaacggctatggaagcaagcctggccttataactacatcaacctcggtcgcagcaaggttggaaggtg  
ggtttatgtgccgaagttcgcgggcgctgcatccacgagtatactggtaacctcgaggctgggtagacaagtagcttactcaagcggctg  
ggtctatctcgaagctccagcttacgacctgccaacgggcagtatggctactccgttgagctattgcgggtgttggtga

SEQ ID NO.: 82

Met Lys Lys Phe Val Ala Leu Phe Ile Thr Met Phe Phe Val Val Ser Met Ala Val Val Ala Gln Pro  
Ala Ser Ala Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val  
Pro Gly Gly Gly Ile Trp Trp Asp Thr Ile Arg Ser Lys Ile Pro Glu Trp Tyr Glu Ala Gly Ile Ser Ala  
Ile Trp Ile Pro Pro Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe  
Phe Asp Leu Gly Glu Tyr Asn Gln Lys Gly Thr Ile Glu Thr Arg Phe Gly Ser Lys Gln Glu Leu Ile  
Asn Met Ile Asn Thr Ala His Ala Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly  
Gly Asp Leu Glu Trp Asn Pro Phe Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly  
Lys Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr Phe  
Gly Gly Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln His Trp Leu Trp Ala Ser Asp Glu Ser  
Tyr Ala Ala Tyr Leu Arg Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala  
Trp Val Val Lys Asp Trp Leu Asn Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val  
Asp Ala Leu Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met  
Asp Glu Ala Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Ser Ala Leu Gln Asn Gly Gln Thr Val Val  
Ser Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr  
Leu Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp  
Leu Asn Lys Asp Arg Leu Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Ser Ile  
Val Tyr Tyr Asp Ser Asp Glu Met Ile Phe Val Arg Asn Gly Tyr Gly Ser Lys Pro Gly Leu Ile Thr  
Tyr Ile Asn Leu Gly Ser Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ala Cys Ile  
His Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Lys Tyr Val Tyr Ser Ser Gly Trp Val Tyr Leu  
Glu Ala Pro Ala Tyr Asp Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

Figure 16 (cont.)

SEQ ID NO.: 83

atggctctggaagacggcggtcataatgcaggccttctactgggatgttctggaggaggaatctggtgggacacaatagctcaaaagata  
cccgaatgggcaagtgcaggaatctcagcgatatggattccaccagcgagtaagggcagtgagcggtggttattccatgggctacgatccctac  
gatttcttgacctggcgagtactatcagaaggggacagttgagacgcgttcggctcaaaaggaagaactggtgaacatgataaacaccgca  
cactctacggcataaaggtgatagcagacatagtcataaaccaccgcgccggtggagacctgagtgaaccccttcgtgaacgactatactt  
ggacagacttctcaaaagtcgctccggtaatatagcccaactaccttgacttccacccaaacgagcttactgttgatgaaggtaaccttg  
gaggataccctgatatgtcacgacaaaagctgggaccagtactggctctgggcgagcagcgaaagctacgctgcctacctcaggagcata  
ggggttgacgctggcggttcgactacgtcaagggctacggagcatgggttgtaacgactggctcagctggtggggaggctgggcccgttgga  
gagtactgggacacgaacgttgatgcactctcaactgggcatacagcagcgccgccaaggtcttgacttcccgtctactacaagatggacg  
aagccttcgacaacaccaacatcccggcattagtgatgcactcagatagcgcagacagtggtcagccgcgacccctcaaggcggttaacttt  
cgttgccaaccacgatacagataatactggaacaagtatccggcttatgcattcatccttacctatgagggacagcctgttatattctaccgcgac  
tacgaggagtggcctaacaaggataagcttaacaacctcatctggatacagatcaccttgctggaggaggagtactgacattgtttactacgacag  
cgacgagcttatcttgtgagaaacggctatggcaccaaccaggactgataacctatatacaacctcggctcaagcaaagttggaaggtgggtc  
tacgttcaaaagtcgccggttcatgcacccagtagtacaccggcaacctcggcggttgatagacaagtacgtctctccagcggctgggtct  
atcttgaggccccagcccagaccggcgacggctactacggctactccgtatggagctactgcggggttggtga

SEQ ID NO.: 84

Met Ala Leu Glu Asp Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly Gly Gly Ile Trp  
Trp Asp Thr Ile Ala Gln Lys Ile Pro Glu Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala  
Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp Leu Gly Glu Tyr  
Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Glu Glu Leu Val Asn Met Ile Asn Thr Ala  
His Ser Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp Leu Glu Trp Asn  
Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn Tyr  
Leu Asp Phe His Pro Asn Glu Leu His Cys Cys Asp Glu Gly Thr Phe Gly Gly Tyr Pro Asp Ile Cys  
His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Ser Glu Ser Tyr Ala Ala Tyr Leu Arg Ser  
Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val Asn Asp Trp Leu  
Ser Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Leu Leu Asn Trp Ala  
Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp Asn Thr  
Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp Pro Phe Lys Ala  
Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile Leu  
Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Lys Leu Asn  
Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr Asp Ser Asp Glu  
Leu Ile Phe Val Arg Asn Gly Tyr Gly Thr Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Gly Ser Ser  
Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr Thr Gly Asn Leu  
Gly Gly Trp Ile Asp Lys Tyr Val Ser Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro Ala His Asp Pro  
Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO.: 85

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cccgaatgggcaagtgcaggaatctcagcgatatggattccaccagcgagtaagggaatgagcggtggttattccatgggctacgatccctac  
gatttcttgacctggcgagtactatcagaaggggacagttgagacgcgttcggctcaaaaggaagaactggtgaacatgataaacaccgca  
cactctacggcataaaggtgatagcggacatagtcataaaccaccgcgccggtggaggcctcagtggaaccccttcgtgaacgactatacc  
tgacagacttctcaaaagtcgctccggtaatatagcccaactaccttgacttccacccaaacgagcttactgttgatgaaggtaaccttg  
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aagccttcgacaataccaacatccccgttggtttacgccctcaagaatggcgggacagtggtcagccgcgacccattcaaggcggttaacttt  
cgttgccaaccacgatacagataatactggaacaagtatccggcttatgcattcatccttacctatgagggacagcctgttatattctaccgcgac  
tacgaggagtggcctaacaaggataagcttaacaacctcatctggatacagatcaccttgctggaggaggagtactgacattgtttactacgacag  
cgacgagcttatcttgtgagaaacggctatggcaccaaccaggactgataacctatatacaacctcggctcaagcaaagctggaaggtgggtc

Figure 16 (cont.)

tacgttccaaagtgcgccgttcacatccacgagtagacacccggcagccctggcggttgatagacaagtacgtctcctccagcggtgggtct  
accttgaggccccggccacgacccggccaatggccagtaggctactccgtctggagctattcgggggtgggtga

SEQ ID NO.: 86

Met Ala Leu Glu Glu Gly Gly Leu Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly Gly Gly Ile Trp  
Trp Asp Thr Ile Ala Gln Lys Ile Pro Glu Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala  
Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp Leu Gly Glu Tyr  
Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Glu Glu Leu Val Asn Met Ile Asn Thr Ala  
His Ser Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Gly Leu Glu Trp Asn  
Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn Tyr  
Leu Asp Phe His Pro Asn Glu Leu His Cys Cys Asp Glu Gly Thr Phe Gly Gly Tyr Pro Asp Ile Cys  
His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Ser Glu Ser Tyr Ala Ala Tyr Leu Arg Ser  
Ile Gly Val Asp Ala Trp Cys Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val Asn Asp Trp Leu  
Ser Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Leu Leu Asn Trp Ala  
Tyr Asn Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp Asn  
Thr Asn Ile Pro Ala Leu Val Tyr Ala Leu Lys Asn Gly Gly Thr Val Val Ser Arg Asp Pro Phe Lys  
Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile  
Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Lys Leu  
Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr Asp Ser Asp  
Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Thr Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Gly Ser  
Ser Lys Ala Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr Thr Gly Ser  
Leu Gly Gly Trp Ile Asp Lys Tyr Val Ser Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro Ala His Asp  
Pro Ala Asn Gly Gln Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO: 87

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gctgccgcccgttacaaaggaacaagccgcagcgacgtaggggtacggagtatacacttgtatgacctggcgaattcaatcaaaaaggga  
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cgaccataaaggcggcgctgacggcacggaatgggtggacgccgtcgaagtcaatccgtccgaccgcaaccaagaaatctcgggcacctat  
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gacgaaagccgaaattgagccgcatttacaaattccggcgcacggcaagcggtgggattgggaagtagacacggaaacggaaactatg  
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gatgggtccggctgatgccgtcaagcatattaagttcagtttttctgattggtgtcgtatgtgcgttctcagactggcaagccgctattaccg  
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gcgcagggattatgcttacggaacgcaacatgattatcttgatcactccgacatcatcggtgggacaagggaagggtcactgaaaaaccagg  
atccgggctggccgcactgatcaccgatgggccgggaggaagcaaatggatgtactgttggaacaacacgctggaaaagtgttctatga

SEQ ID NO: 88

Met Phe Leu Leu Ala Phe Leu Leu Thr Ala Ser Leu Phe Cys Pro Thr Gly Gln Pro Ala Lys Ala Ala  
Ala Pro Phe Asn Gly Thr Met Met Gln Tyr Phe Glu Trp Tyr Leu Pro Asp Asp Gly Thr Leu Trp Thr  
Lys Val Ala Asn Glu Ala Asn Asn Leu Ser Ser Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro Ala Tyr  
Lys Gly Thr Ser Arg Ser Asp Val Gly Tyr Gly Val Tyr Asp Leu Tyr Asp Leu Gly Glu Phe Asn Gln  
Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr Lys Ala Gln Tyr Leu Gln Ala Ile Gln Ala Ala His Ala  
Ala Gly Met Gln Val Tyr Ala Asp Val Val Phe Asp His Lys Gly Gly Ala Asp Gly Thr Glu Trp Val  
Asp Ala Val Glu Val Asn Pro Ser Asp Arg Asn Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp  
Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr His Phe Asp

Gly Val Asp Trp Asp Glu Ser Arg Lys Leu Ser Arg Ile Tyr Lys Phe Arg Gly Ile Gly Lys Ala Trp  
Asp Trp Glu Val Asp Thr Glu Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp  
His Pro Glu Val Val Thr Glu Leu Lys Asn Trp Gly Lys Trp Tyr Val Asn Thr Thr Asn Ile Asp Gly  
Phe Arg Leu Asp Ala Val Lys His Ile Lys Phe Ser Phe Phe Pro Asp Trp Leu Ser Tyr Val Arg Ser  
Gln Thr Gly Lys Pro Leu Phe Thr Val Gly Glu Tyr Trp Ser Tyr Asp Ile Asn Lys Leu His Asn Tyr  
Ile Thr Lys Thr Asp Gly Thr Met Ser Leu Phe Asp Ala Pro Leu His Asn Lys Phe Tyr Thr Ala Ser  
Lys Ser Gly Gly Ala Phe Asp Met Arg Thr Leu Met Thr Asn Thr Leu Met Lys Asp Gln Pro Thr  
Leu Ala Val Thr Phe Val Asp Asn His Asp Thr Glu Pro Gly Gln Ala Leu Gln Ser Trp Val Asp Pro  
Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly  
Asp Tyr Tyr Gly Ile Pro Gln Tyr Asn Ile Pro Ser Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile Ala Arg  
Arg Asp Tyr Ala Tyr Gly Thr Gln His Asp Tyr Leu Asp His Ser Asp Ile Ile Gly Trp Thr Arg Glu  
Gly Val Thr Glu Lys Pro Gly Ser Gly Leu Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser Lys Trp  
Met Tyr Cys Trp Gln Thr Thr Arg Trp Lys Ser Val Leu

atgaaagaagcgggtgtgtatcaaattttcccggtatcggttcttaattggcaaccttcaaatgataacagcaagcagcaggcacgcggggcgccgagccgattgagcatcgcgatttggcgtgatttggccgataatccgcgcctgaaaggggacgagcggctacgatggcgacgggtgaatggcgaatgactttttcggcgagacatcgcgggaattgaacaaaagtggattatttgcagtcgcttggagtgaaacacgattacttaaatccgatcgccaatgcgccatcgaaccataaataatgatgcgagcaattacaagaattggatccgatgttcggttccccggaagaattccaatcggttgtgcaggcgcttgcgaaccgggggatgcatctcatcttagacgggggtgttcaaccacgtatccgacgattcgatttaccgctaccaccgctatccgaccgtcgggtgcgtatgaatatggggaagcggtttacgatttgaatgaaaggaattgagcggaggaagaagcgcggaaacaagtggagagaaagttcaacaagaggggacagacgttcagcccgtatgggtttcatcttgggttcaatattgaaaacaaaaagtcattggccattatcaataccaatcatggtggggctatgacagctcgcggagtttaagtcggtgacgggggaaaaagtgcgcacccgagtgaaatgaacaacgatgcgctcgcgaattacattttccgtgaatcggaattcgggtggcgaaaaagctggattgccctcggcgccctcggctggcggttggatgtggccaatgaggttgatccggcgctttggcgcgagtttcccaagaattgcttcaagggtcgtacggccgcgggtccgacgttaaaagagggggagcagccgctcatttaggggaaatttgggatgacgcacgaaatattttctaggcgaccagtagattccggtgatgaactaccgggtccgcggggcggtgcttgacttttgaaaaacggaaatgcagaagaggcgggacaagcggctgacggccataagggaagactacccaagtgaagcgtttatgcgctgatgaacttaatcggttcgcatgacacggcgcgggcggtcttctgcttgggaacggaacggattacccgagcggcggaagcttgatccgaattataatgaggaacttgggaaaaagcgggctcaagctggcggtgattttgcagatgggataccggggagcgcgcgacgatttattacggcgatgaagcgggagtaacaggctcaaaagacccagacaaccgccgcacgtatccgtggggcaagaagatcaaaatctgttgcctattatcagaagtggggacattcgccagcaccatcaatcggttgggttggccatggcgacatacaagacgggtgatgcgcaagggtatgtatacgtatttcccgccaatcggggcggtgaagcggcgtcattgccatcaaccgcggcaatgaggacaagacgggtggcgcttgacgtcgcttcgttccgaacggcaccgtgcttacggatgagttgcatgatggcggggaagctacggctcgtggcggaacgttgacgggtcacgattccggccctggatggacggatgatgttgggacgggtgacggcggaatgccggcagcagtcagcaatttgcaggcgagcgttcggatggctgcgtgacgttaacgtgggaaggaatgcatcgagataccgaattttagagttccagcttaaaaagggtgccggttatacgatggtgcaagagacggaaacaacttcggccacgatcggttcgttgacgaacggaaacgcctattacttgcggttgcggcggtcgtgatgaaaacgggaatgaatcacccaaggtcgaaacgaatcgctcgttcctcattaccgcgtgacgagcgacaatgtccagttcgtgacaacgttaagcgtatccacactggatttgcaaaagccgcagcaagtggatgtccatgtcaacatcgacaatgtgacaagcaaaaggagcagctgatgggttgcaagcgggtgttgcaagtgaaaggccgcatgacgaaacatggaaagaatcacagcggcttaccaggacaagacggcgacgccaacgtgttcgagctgccttcaactccgctcgcgcgaggagctatacgtatcgttatgcgtgacgaccaaccttggcgaggaagtggatgtatacagaagagaagcaagtgcgttgcggcgagacaacagcgaccaaatagcgccagcagacgccatcagctgcggcgagcctgcgggtgaatcgggacaagtgaatttatcatggacgttgttgggaaaaagatggggatgcttattgttagccatcgagcgcaacgggtgatcgtgcatacaaccacttcgatcggcgattcattacagactacgatgtcgaaaacggcaccgagtagacgtatgtgtgcaaggttgatgaccgcgggcaatgtgtggcgtcaaacacgggtcaaggtgacggcggacattgtgatggtgaaagtattttaaagtgagagcgccggattacacaccgttggatgccgaattacgattccgaacagcttgaacggctggaacacagggggcctgggagatgtcgcgcaacgggtcgggtgacggccgattggcaatttaccgtcgaggtgcaggaaggggaaacgatacctaataagtatgtgaaaggcggtatcgtggatcaagagggttggccgaccatacgcgtgaggacgacaacgatgatgacgtgagctactacggctatgggacgattggcaccgacttgaagtgcggtcacaaatgaaggaaacaatcagatgattgtgcaagaccgattttgcgtggatcgatatgccggtcgtatcgaagaggtgcaaaaacaagggaagtcaagtgcagatcaagggaatgccattaaaaacgggtgtttgacgatcaatggcgagcgggtgccgattgatggccgatggttcctgtacacggttgcggcgccagccatcaaaaaagaagtgttgatccatatgaaccatcggccgaaagcaaaaacgccattttcaaacgacggcg

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
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gcccacgattttccacccgggtggggtaccaggta

Met Lys Glu Ala Val Val Tyr Gln Ile Phe Pro Asp Arg Phe Phe Asn Gly Asn Ser Ser Asn Asn Ser Lys Gln Gln Ala Arg Gly Ala Gln Pro Ile Glu His Arg Asp Trp Ser Asp Leu Pro Asp Asn Pro Arg Leu Lys Gly Thr Ser Gly Tyr Asp Gly Asp Gly Glu Trp Ser Asn Asp Phe Phe Gly Gly Asp Ile Ala Gly Ile Glu Gln Lys Leu Asp Tyr Leu Gln Ser Leu Gly Val Asn Thr Ile Tyr Leu Asn Pro Ile Ala Asn Ala Pro Ser Asn His Lys Tyr Asp Ala Ser Asn Tyr Lys Glu Leu Asp Pro Met Phe Gly Ser Pro Glu Glu Phe Gln Ser Phe Val Gln Ala Leu Ala Asn Arg Gly Met His Leu Ile Leu Asp Gly Val Phe Asn His Val Ser Asp Asp Ser Ile Tyr Phe Asp Arg Tyr His Arg Tyr Pro Thr Val Gly Ala Tyr Glu Tyr Trp Glu Ala Val Tyr Asp Leu Met Asn Glu Lys Gly Leu Ser Glu Glu Glu Ala Arg Lys Gln Val Glu Glu Lys Phe Lys Gln Glu Gly Gln Thr Phe Ser Pro Tyr Gly Phe His Leu Trp Phe Asn Ile Glu Asn Lys Lys Val Asn Gly His Tyr Gln Tyr Gln Ser Trp Trp Gly Tyr Asp Ser Leu Pro Glu Phe Lys Ser Val Thr Gly Glu Lys Val Pro His Pro Ser Glu Leu Asn Asn Asp Ala Leu Ala Asn Tyr Ile Phe Arg Glu Ser Asp Ser Val Ala Lys Ser Trp Ile Ala Leu Gly Ala Ser Gly Trp Arg Leu Asp Val Ala Asn Glu Val Asp Pro Ala Phe Trp Arg Glu Phe Arg Gln Glu Leu Leu Gln Gly Ser Tyr Gly Arg Gly Pro Thr Leu Lys Glu Gly Glu Gln Pro Leu Ile Leu Gly Glu Ile Trp Asp Asp Ala Ser Lys Tyr Phe Leu Gly Asp Gln Tyr Asp Ser Val Met Asn Tyr Arg Phe Arg Gly Ala Val Leu Asp Phe Leu Lys Asn Gly Asn Ala Glu Glu Ala Asp Lys Arg Leu Thr Ala Ile Arg Glu Asp Tyr Pro Ser Glu Ala Phe Tyr Ala Leu Met Asn Leu Ile Gly Ser His Asp Thr Ala Arg Ala Val Phe Leu Leu Gly Asn Gly Thr Asp Ser Ser Glu Arg Ala Glu Leu Asp Pro Asn Tyr Asn Glu Glu Leu Gly Lys Lys Arg Leu Lys Leu Ala Val Ile Leu Gln Met Gly Tyr Pro Gly Ala Pro Thr Ile Tyr Tyr Gly Asp Glu Ala Gly Val Thr Gly Ser Lys Asp Pro Asp Asn Arg Arg Thr Tyr Pro Trp Gly Lys Glu Asp Gln Asn Leu Leu Ser His Tyr Gln Lys Val Gly His Ile Arg Gln His His Gln Ser Leu Leu Ala His Gly Asp Ile Lys Thr Val Tyr Ala Gln Gly Asp Val Tyr Val Phe Ala Arg Gln Tyr Gly Arg Glu Ala Ala Leu Ile Ala Ile Asn Arg Gly Asn Glu Asp Lys Thr Val Ala Leu Asp Val Ala Ser Leu Leu Pro Asn Gly Thr Val Leu Thr Asp Glu Leu His Asp Gly Gly Glu Ala Thr Val Ala Gly Gly Thr Leu Thr Val Thr Ile Pro Ala Leu Asp Gly Arg Met Met Phe Gly Thr Val Thr Ala Glu Met Pro Ala Ala Val Ser Asn Leu Gln Ala Ser Ala Ser Asp Gly Cys Val Thr Leu Thr Trp Glu Gly Asn Ala Ser Arg Tyr Arg Ile Tyr Glu Ser Thr Leu Lys Gly Ala Gly Tyr Thr Met Val Gln Glu Thr Glu Thr Thr Ser Ala Thr Ile Gly Ser Leu Thr Asn Gly Thr Ala Tyr Tyr Phe Ala Val Ala Ala Val Asp Glu Asn Gly Asn Glu Ser Pro Lys Val Glu Thr Asn Arg Val Val Pro His Tyr Pro Leu Thr Ser Asp Asn Val Gln Phe Val Thr Thr Leu Ser Asp Ala Thr Leu Asp Leu Ser Lys Pro Gln Gln Val Asp Val His Val Asn Ile Asp Asn Val Thr Ser Lys Gly Ala Ala Asp Gly Leu Gln Ala Val Leu Gln Val Lys Gly Pro His Asp Glu Thr Trp Lys Glu Tyr Arg Ala Ala Tyr Gln Gly Gln Asp Gly Asp Ala Asn Val Phe Arg Ala Ala Phe Thr Pro Leu Ala Ala Gly Thr Tyr Thr Tyr Arg Tyr Ala Leu Thr Thr Asn Leu Gly Glu Glu Trp Met Tyr Thr Glu Glu Lys Gln Val Thr Phe Ala Ala Asp Asn Ser Asp Gln Ile Ala Pro Ala Asp Ala Ile Glu Leu Arg Gln Pro Ala Val Glu Ser Gly Gln Val Asn Leu Ser Trp Thr Phe Val Gly Lys Lys Asp Gly Asp Ala Tyr Leu Leu Ala Ile Glu Arg Asn Gly Asp Ile Val His Thr Thr Thr Ser Ile Gly Asp Ser Phe Thr Asp Tyr Asp Val Glu Asn Gly Thr Glu Tyr Thr Tyr Val Val Lys Leu Tyr Asp Arg Ala Gly Asn Val Val Ala Ser Asn Thr Val Lys Val Thr Pro Asp Ile Val Met Val Lys Val Ile Phe Lys Val Arg Ala Pro Asp Tyr Thr Pro Leu Asp Ala Arg Ile Thr Ile Pro Asn Ser Leu Asn Gly Trp Asn Thr Gly Ala Trp Glu Met Ser Arg Asn Gly Ala Val Thr Pro Asp Trp Gln Phe Thr Val Glu Val Gln Glu Gly Glu Thr Ile Thr Tyr Lys Tyr Val Lys Gly Gly Ser Trp Asp Gln Glu Gly Leu Ala Asp His Thr Arg Glu Asp Asp Asn Asp Asp Asp Val Ser Tyr Tyr Gly Tyr Gly Thr Ile Gly Thr Asp Leu Lys Val Thr Val His Asn Glu Gly Asn Asn Thr Met Ile Val Gln Asp Arg Ile Leu Arg Trp Ile Asp Met Pro Val Val Ile Glu Glu Val Gln Lys Gln Gly Ser Gln Val Thr Ile Lys Gly Asn Ala Ile Lys Asn Gly Val Leu Thr Ile Asn Gly Glu Arg Val Pro Ile Asp Gly Arg Met Ala Phe Ser Tyr Thr Phe Ala Pro Ala Ser His Gln Lys Glu Val Leu Ile His Ile Glu Pro Ser Ala Glu Ser Lys Thr Ala Ile Phe Asn Asn Asp

Figure 16 (cont.)

Gly Gly Ala Ile Ala Lys Asn Thr Lys Asp Tyr Val Leu Asn Leu Glu Thr Lys Gln Phe Lys Lys Leu  
Leu Glu Ser Thr Ser Arg Ala Ala Ala Gly Pro Ser Ile Phe His Pro Gly Gly Val Pro Gly

SEQ ID NO: 91

gtgctaacgtttaccgcatcattegaaaaggatggatgttctgctcgcgtttttgtcactgectcgtgttctgcccaacaggacagcccgcca  
aggctgccgcaccgtttaacggcaccatgatgcagtatgttgatggtactgcccggatgatggcacgttatggaccaaagtggccaatgaagc  
caacaactatccagccttggcatcaccgctctttggctgccgcccgttataaaggaaacagccgcagcgacgtagggtagcgagatacga  
cttgatgacctcggcgaattcaatcaaaaaggacgctccgcacaaaatacggaaacaaagctcaatatcttcaagccattcaagccgccac  
gccgctggaatgcaagtgtacgccgatgctgtgttcgaccataaaggcggcgccgacggcacggaatgggtggacgccgctgaagtcattc  
cgctcgaccgcaaccaagaaatcctgggcacctatcaaatccaagcatggacgaaatttgatttccggggcggggcaacacctactccagctt  
taagtggcgtgtgtaccattttgacggcgttgattgggacgaaagccgaaattgagccgcatttacaaattccggcgcatcggaagcgtgg  
gattgggaagtagacacggaaaacggaaactatgactacttaattgatgccgacttgacatggaccatcctgaagtgtgtacggaactgaaaa  
actggggcaaatggtatgtcaacacacgaacattgatgggttcgcttgatgccgtcaagcatattaaagttcagtttttctgattggtgtcgt  
atgtcgttctcagactggcaagccgctattaccgctggggaatttgagctatgacatcaacaagttgcacaattacattacgaaaacaaag  
gaacgatgtctttgttgatgccccgttacacaacaaatttataccgcttccaaatcaggggggcgcatgttgatgacgcacgttaataaccaatact  
ctcatgaaagatcaaccgacattggccgctaccttcgttgataatcatgacaccgaaccggccaagcgtgcagtcattgggtcgacccatggt  
tcaaaccgttggttacgctttattctaacctggcaggaaggataccgctgcgtctttatggtgactattatggcatcccaataataacattcctt  
cgctgaaaagcaaatcgatccgctcctcatcgccgcagggattatgcttacggaacgcaacatgattatcttgatcactccgacatcatcggt  
tggacaagggaaggcgtcactgaaaaccaggatccggactggccgactgatcaccgatgggcccgggaggaagcaaatggatgtacgtt  
ggcaaacacacgccggaaggtgtctatgacctaccggcaaccggagtgacaccgtcaccatcaacagtgatggatggggagaattcaa  
agtcaatggcggcttcggttgggttgcctagaaaaacgaccgctctaccatcgcttggccgatcacaacccgaccgtggactggtgaatt  
cgctcgttggaacgaaccacggttggtggcatggccttga

SEQ ID NO: 92

Val Leu Thr Phe His Arg Ile Ile Arg Lys Gly Trp Met Phe Leu Leu Ala Phe Leu Leu Thr Ala Ser  
Leu Phe Cys Pro Thr Gly Gln Pro Ala Lys Ala Ala Ala Pro Phe Asn Gly Thr Met Met Gln Tyr Phe  
Glu Trp Tyr Leu Pro Asp Asp Gly Thr Leu Trp Thr Lys Val Ala Asn Glu Ala Asn Asn Leu Ser Ser  
Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys Gly Thr Ser Arg Ser Asp Val Gly Tyr Gly  
Val Tyr Asp Leu Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr  
Lys Ala Gln Tyr Leu Gln Ala Ile Gln Ala Ala His Ala Ala Gly Met Gln Val Tyr Ala Asp Val Val  
Phe Asp His Lys Gly Gly Ala Asp Gly Thr Glu Trp Val Asp Ala Val Glu Val Asn Pro Ser Asp Arg  
Asn Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn  
Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr His Phe Asp Gly Val Asp Trp Asp Glu Ser Arg Lys Leu  
Ser Arg Ile Tyr Lys Phe Arg Gly Ile Gly Lys Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn  
Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp His Pro Glu Val Val Thr Glu Leu Lys Asn  
Trp Gly Lys Trp Tyr Val Asn Thr Thr Asn Ile Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys  
Phe Ser Phe Phe Pro Asp Trp Leu Ser Tyr Val Arg Ser Gln Thr Gly Lys Pro Leu Phe Thr Val Gly  
Glu Tyr Trp Ser Tyr Asp Ile Asn Lys Leu His Asn Tyr Ile Thr Lys Thr Asn Gly Thr Met Ser Leu  
Phe Asp Ala Pro Leu His Asn Lys Phe Tyr Thr Ala Ser Lys Ser Gly Gly Ala Phe Asp Met Arg Thr  
Leu Met Thr Asn Thr Leu Met Lys Asp Gln Pro Thr Leu Ala Val Thr Phe Val Asp Asn His Asp  
Thr Glu Pro Gly Gln Ala Leu Gln Ser Trp Val Asp Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile  
Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly Asp Tyr Tyr Gly Ile Pro Gln Tyr Asn Ile  
Pro Ser Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln His  
Asp Tyr Leu Asp His Ser Asp Ile Ile Gly Trp Thr Arg Glu Gly Val Thr Glu Lys Pro Gly Ser Gly  
Leu Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln His Ala Gly  
Lys Val Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn Ser Asp Gly Trp Gly Glu  
Phe Lys Val Asn Gly Gly Ser Val Ser Val Trp Val Pro Arg Lys Thr Thr Val Ser Thr Ile Ala Trp  
Pro Ile Thr Thr Arg Pro Trp Thr Gly Glu Phe Val Arg Trp Thr Glu Pro Arg Leu Val Ala Trp Pro

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Figure 16 (cont.)

SEQ ID NO: 93

atgaaatcgtttgcattcatgcctatcctttttatgcaaacgatttcatcagtgaaagggaaggaggagaaaaatggggaagaatatgagaaga  
agattcacgtatttttcaatcttcttattgttcgttcagctgtttcatttagtgaaccgctagcgccaatggaacggtgaacagtagtctgtgtgtaa  
tggaacagagtcacgtttctatatggaggaacaggaacgagcagctctgttactggcaggctcctttaatgattggcagaaagatggtgaca  
agaagattgcactaacaagaagcgcacaataacgtctgtgttcacgcaaacacttcaagatgggacataacgtataagtttgtgtagatggtc  
aatgggtggcggatccgcttaacccgaatcaagtagacgacggttacggcgccgtaatagtgtcgttgttgcggacaccggtgcaacaag  
aacggacagtgacgcttgttgtaacttacaagacgaattaggtcatacagcggaatgggatccgaaagcgacagctacgtatgaaaaagg  
aagggaacgggttatatacgtttacaggtacactccagccggaacgtacagatataaaattgcgattaatggcagctgggacgaaaactatggt  
gtcggcgccgcgatggcggaatattaagctgtattaaatgaacaacaacggttacattttattacaacgacagaacgcattgcgattgcgg  
attcgacttggtagcaccaattctaaaagaaaagcagccgcggctcgttgaacgattttaccagctattggttatgaaacagacgtgaacggtt  
ggacgccgcaaacatcaacggcgttgttgcagatgatttgaattcatttatacgtttaaaggcgcgtgtgccaaaaggacatatgaataaa  
agtagttcttgggaatgattggacatatgaaaattatccacaagataatgccaaattaaatgtgttgaagaacgacaattaccttttcttaacgc  
gaaaacgaaagtagtgataaccgattacaatccaagcgggtcggatggtatcgtcaaaaagaccgttgaagcataatcgtgggattcgttga  
tcgccaaccgttgggtgcgggtgaaagctgggacagaagtgacccttcgtttatcagcgaaaaagggtgattgacaaaagcggtatgtatgtaa  
aaaatacgacaaccggcacagcgaaactatattcgatgaaaaagccgggtgttcttggcgaagaagaatttgggaagcgacattcacaccgg  
atgtgaaaggagtatacgggtataaattattgcggtagatgttggaacgaaagcagaatacggggaagatacacaagaagggcagtgggga  
aaagcagtagataaaaatgcagagctgttccaattaacggtgtacgacctcctaccaaacaccggattggtgaaagaagcagttgtatatca  
aattttccctgatccaaag

SEQ ID NO: 94

Met Lys Ser Phe Ala Phe Met Pro Ile Leu Phe Tyr Ala Asn Asp Phe Ile Ser Glu Arg Glu Gly Gly  
Gly Lys Met Gly Lys Asn Met Arg Arg Arg Phe Thr Tyr Phe Ser Ile Phe Leu Leu Phe Val Gln Leu  
Phe Ser Phe Ser Ala Thr Ala Ser Ala Asn Gly Thr Val Asn Ser Ser Pro Val Val Asn Gly Asn Glu  
Val Thr Phe Leu Tyr Gly Gly Thr Gly Asn Glu Gln Ser Val Leu Leu Ala Gly Ser Phe Asn Asp Trp  
Gln Lys Asp Gly Asp Lys Lys Ile Ala Leu Thr Lys Gly Asp Asn Asn Val Trp Ser Val Thr Gln Thr  
Leu Gln Asp Gly Thr Tyr Thr Tyr Lys Phe Val Val Asp Gly Gln Trp Val Ala Asp Pro Leu Asn Pro  
Asn Gln Val Asp Asp Gly Tyr Gly Gly Arg Asn Ser Val Val Val Val Gly Thr Pro Val Gln Gln Glu  
Arg Thr Val Thr Leu Val Gly Asn Leu Gln Asp Glu Leu Gly His Thr Ser Glu Trp Asp Pro Lys Ala  
Thr Ala Thr Val Met Lys Lys Glu Gly Asn Gly Leu Tyr Thr Phe Thr Gly Thr Leu Pro Ala Gly Thr  
Tyr Glu Tyr Lys Ile Ala Ile Asn Gly Ser Trp Asp Glu Asn Tyr Gly Val Gly Gly Arg Asp Gly Gly  
Asn Ile Lys Leu Leu Leu Asn Glu Gln Thr Thr Val Thr Phe Tyr Tyr Asn Asp Arg Thr His Ala Ile  
Ala Asp Ser Thr Trp Tyr Ala Pro Ile Leu Lys Glu Lys Gln Pro Arg Leu Val Gly Thr Ile Leu Pro  
Ala Ile Gly Tyr Glu Thr Asp Val Asn Gly Trp Thr Pro Gln Thr Ser Thr Ala Leu Leu Ser Asp Asp  
Asp Phe Asp Ser Ile Tyr Thr Phe Lys Ala Arg Val Pro Lys Gly Thr Tyr Glu Tyr Lys Val Val Leu  
Gly Asn Asp Trp Thr Tyr Glu Asn Tyr Pro Gln Asp Asn Ala Lys Leu Asn Val Leu Glu Glu Thr  
Thr Ile Thr Phe Phe Phe Asn Ala Lys Thr Lys Val Val Tyr Thr Asp Tyr Asn Pro Ser Gly Ser Asp  
Gly Ile Val Gln Lys Asp Arg Leu Lys His Asn Thr Trp Asp Ser Leu Tyr Arg Gln Pro Phe Gly Ala  
Val Lys Ala Gly Thr Glu Val Thr Leu Arg Leu Ser Ala Lys Lys Gly Asp Leu Thr Lys Ala Asp Val  
Tyr Val Lys Asn Thr Thr Thr Gly Thr Ala Lys Leu Tyr Ser Met Lys Lys Ala Gly Val Leu Gly Glu  
Glu Glu Tyr Trp Glu Ala Thr Phe Thr Pro Asp Val Lys Gly Val Tyr Gly Tyr Lys Phe Ile Ala Val  
Asp Ala Gly Thr Lys Ala Glu Tyr Gly Glu Asp Thr Gln Glu Gly Gln Trp Gly Lys Ala Val Asp Lys  
Asn Ala Glu Leu Phe Gln Leu Thr Val Tyr Asp Pro Ser Tyr Gln Thr Pro Asp Trp Met Lys Glu Ala  
Val Val Tyr Gln Ile Phe Pro Asp Pro Lys

SEQ ID NO: 95

atgtatacactattcatccgttcataatttggatactgatggtgatggtgtaggagactttagtgagttgctgaaaaggtagattatctaaatctcttg  
gagtagatacagctcgtgttttaccatttaataaaagtaaaacttatcatggatatgatgttgaagattactatgatgtagaaccagattatggaacact  
acaagatcttgataatatgataaaagtctaaatgaaaatggaataaaggtagtaatggatcttgttgaatcatacgtcggatacacatccatggtt  
tcttgatgcagttgaaaatactactaatttcctatattggaactattacattatgagcttggatgagcctcaaaataagaatcattggcattataaggtt  
aattcaaaaggacaaactgtgtggtattttggattgttgattcatcaatccggacctaattacgacaaccctaagtaatggtgaagtgaataa

Figure 16 (cont.)

aataatagattttgggcagatatgggagtagatggatttagattagatgcagcaaacattattatggattgactggagcgatggaattgaacag  
tcagcaagcgttgcacaaagagatagaagactatataaaagataaaactaggggaaaatgcaatagttgtgagtgagggttacgatggagattcaa  
atgttcttttaaaatttgcctcaatgcctgtgttaatttttagttttgtacaatttgagaggaaatttgaaggagagataactaatttcagactctatt  
agttgggttgattcctcgtgtataatttaaatgttttccattttgtagatcatgacgttgacagattatttctgagctttagatagtaaatatc  
aggagatgtaatactgccacaaaacaattttgctagttaattgcttactactctcattaacaggcatgccaaactatttactatggtgatgaaatag  
gacttaggggagtggaagtggcattcagaacctgggatataacctgtgcgtgagccaatgcaatggataaggatcaaaaagggaacggcaca  
cttattggacaaaagagttttacgaaggtattactgaaggagtctaatgaagatggagcaatatacatgatccagatgatggagtatctgtag  
aagaacaagaaaatggatattctattttaaaactttttaagaatttatcaacttacgaaaagattatccggcacttgcctttggaagtactacgattga  
gagagattggaaaaactgtatgttttgaaaaagtcgtataacttccaggatgttctgtattaattaacctgtatccaacgtattcaatacatacgaa  
gtccagaagggtataaatgggtgtggtatgcattttttagtggtgacaactatgaatttggagcaaaaagatgaaatgattttacagaatacaagt  
gacgataaatccaaggcaaatatttatattgttaaagtaa

SEQ ID NO: 96

Met Tyr Thr Leu Phe Ile Arg Ser Tyr Phe Asp Thr Asp Gly Asp Gly Val Gly Asp Phe Ser Gly Val  
Ala Glu Lys Val Asp Tyr Leu Lys Ser Leu Gly Val Asp Thr Val Trp Phe Leu Pro Phe Asn Lys Ser  
Lys Ser Tyr His Gly Tyr Asp Val Glu Asp Tyr Tyr Asp Val Glu Pro Asp Tyr Gly Thr Leu Gln Asp  
Leu Asp Asn Met Ile Lys Val Leu Asn Glu Asn Gly Ile Lys Val Val Met Asp Leu Val Val Asn His  
Thr Ser Asp Thr His Pro Trp Phe Leu Asp Ala Val Glu Asn Thr Thr Asn Ser Pro Tyr Trp Asn Tyr  
Tyr Ile Met Ser Leu Asp Glu Pro Gln Asn Lys Asn His Trp His Tyr Lys Val Asn Ser Lys Gly Gln  
Thr Val Trp Tyr Phe Gly Leu Phe Asp Ser Ser Met Pro Asp Leu Asn Tyr Asp Asn Pro Lys Val Met  
Asp Glu Val Lys Lys Ile Ile Asp Phe Trp Ala Asp Met Gly Val Asp Gly Phe Arg Leu Asp Ala Ala  
Lys His Tyr Tyr Gly Phe Asp Trp Ser Asp Gly Ile Glu Gln Ser Ala Ser Val Ala Lys Glu Ile Glu  
Asp Tyr Ile Lys Asp Lys Leu Gly Glu Asn Ala Ile Val Val Ser Glu Val Tyr Asp Gly Asp Ser Asn  
Val Leu Leu Lys Phe Ala Pro Met Pro Val Phe Asn Phe Ser Phe Met Tyr Asn Leu Arg Gly Asn  
Phe Glu Gly Arg Asp Asn Leu Ile Ser Asp Ser Ile Ser Trp Val Asp Ser Ser Leu Tyr Asn Leu Asn  
Val Phe His Phe Pro Phe Ile Asp Ser His Asp Leu Asp Arg Phe Ile Ser Glu Leu Val Asp Ser Lys  
Tyr Gln Gly Asp Val Ile Ser Ala Thr Lys Gln Tyr Leu Leu Val Asn Ala Leu Leu Leu Ser Leu Thr  
Gly Met Pro Thr Ile Tyr Tyr Gly Asp Glu Ile Gly Leu Arg Gly Trp Lys Trp His Ser Glu Pro Trp  
Asp Ile Pro Val Arg Glu Pro Met Gln Trp Tyr Lys Asp Gln Lys Gly Asn Gly Gln Thr Tyr Trp Thr  
Lys Glu Phe Tyr Glu Gly Ile Thr Glu Gly Ser Ala Asn Glu Asp Gly Ala Ile Tyr Asp Asp Pro Asp  
Asp Gly Val Ser Val Glu Glu Gln Glu Asn Gly Tyr Ser Ile Leu Asn Phe Phe Lys Glu Phe Ile Asn  
Leu Arg Lys Asp Tyr Pro Ala Leu Ala Phe Gly Ser Thr Thr Ile Glu Arg Asp Trp Lys Asn Leu Tyr  
Val Leu Lys Lys Ser Tyr Asn Phe Gln Asp Val Leu Val Leu Ile Asn Leu Asp Pro Thr Tyr Ser Asn  
Thr Tyr Glu Val Pro Glu Gly Tyr Lys Trp Val Trp Tyr Ala Phe Phe Asp Gly Asp Asn Tyr Glu Phe  
Gly Ala Lys Asp Glu Met Ile Leu Gln Asn Thr Ser Trp Thr Ile Asn Pro Arg Gln Ile Tyr Ile Phe Val  
Lys

SEQ ID NO: 97

atgaggaagaagatgtcgcattcaagattacttttctttgatcttagcactttttatttcttccggttgatttcagaagttaaaagcgaagccag  
ctactaaattcaaaagcaaaaggtcctgtaaaagtaaatgttaatacgccatttattgagaatgctactactaatacgtggagtgttcaaaagaatc  
tttattgattatcttagtaaagtattactgttaaggatgtaaatgatcagattgtattactaaggaaacaacgaacaaaacaatatttttgaa  
attgaacttctcctggaacttatacatttgaggtaaaaggatatgaggaagatttagttatatttccaggggaaaaagttaatcagatcatagatgag  
aaaaataatattgtaattgtcgaactttttgttaagtgaatagttaggacaataatgaagttgacgatattttataaaaaattatgatattacatcgg  
caacgttgatcttcaaaaaagatacagcacaagaagattatgaagggtacctgtaaccttacaggtacttccactttaattaataaagaattatat  
cctggtatgtggactgtaaaatttgaagttgatcttaaatcaaggatgcaagatgttaccagaaaaagttcatcttgaaaaatgaatttagcataga  
agtgtccagcaaaagacaaaaagttaacatttaagtgttagtctttgatacagaggttaatagaaccgaaattagtagttgtatttccgcaaatgaggt  
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caagaattcgataaattatgtgaatcgctattaatgtttatgccaacggtaaagagagtggttagttgttcaaaaaagaaaattataaacttata  
gatttagaaagtgttgacagtataagtgtacttataacgttgatagcatgaatgagcttaagttggattggaattatccaattcaagttgacttttgaag

Figure 16 (cont.)

tttgaaaaaaggtataaatagcaatgaatacgaataatcttcaactaacacaaaattcttttcaacagaattcacaggcaggcaatttgggatac  
ttgagaaaaattgcgattagtagtgctaatggatttgaagtaagattaatgagatttcaagagatgataaactataacatcattgaatcttctct  
tacatcgtctactatgtatacactattcatccgttcataatgatactgatgggtgatggtaggagactttagtggagttgctgaaaaggtagattatc  
taaatctcttggagtagatacagctcgtgttttaccatttaataaaaagtaaatcttatcatggatatgatgttgaagattactatgatgagaaccagat  
tatggaacactacaagatcttgataatgataaaaagtctaaatgaaaatggaataaaggtagtaattggatcttgttgaatcatacgtcggatac  
acatccatggtttcttgatgcagttgaaaatactactaattctccatattggaactattacattatgagcttggatgagcctcaaaataagaatcattgg  
cattataagggttaattcaaaaaggacaaactgtgtggtattttggattgtttgattcatcaatgccggacctaattacgacaacctaaagtaattggat  
gaagtgaaaaaataatagatttttgggcagatatgggagtagatggatttagattagatgcagcaaacattattatggatttactggagcgtatg  
gaattgaacagtcagcaagcgttgcacaaagagatagaagactatataaaagataaaactaggggaaaaatgcaatagtgtgagtgaggtttacga  
tggagattcaaatgttcttttaaaattgtcctcaatgcctgtgttaatttttagttttatgtacaatttgagaggaaatttgaaggagagataacttaatt  
tcagactctattagtgtgggttattcctcgtgtataatttaaatgttttccattttcattttgatatgtagcatgcttgacagatttttctgagctttag  
atagtaaatatcagggagatgtaatatctgccacaaaacaatatttgcagtaaatgtcttactactctcattaacaggcatccaactatttactatgg  
tgatgaaataggacttaggggatggaagtggcattcagaacctgggataacctgtgcgtgagccaatgcaatggtataaggatcaaaaagg  
gaacggtcaaacattattggacaaaagagttttacgaagggtattactgaagggaagtgtcaatgaagatggagcaatatacagatccagatgatg  
gagtagtctgtagaagaacaagaaaatggatatcttatttttaaaacttttttaagaatttatcaacttacgaaaagattatccggcacttgcctttggaagt  
actacgattgagagagattggaaaaactgtatgtttgaaaaagtcgtataactccaggatgttctgtattaattaaccttgatccaacgtattcaa  
atacatcgaagttccagaagggtataaatgggtgtggtatgcatttttgatggtgacaactatgaatttggagcaaaagatgaaatgattttacg  
aatacaagttggacgataaatccaaggcaaatatttatatttgaagtaa

SEQ ID NO: 98

Met Arg Lys Lys Met Ser His Ser Arg Phe Thr Phe Leu Leu Ile Leu Ala Leu Phe Ile Phe Phe Ser  
Gly Cys Ile Ser Glu Val Lys Ser Glu Ser Gln Leu Leu Asn Ser Lys Gln Lys Val Leu Val Lys Val  
Asn Val Asn Thr Pro Phe Ile Glu Asn Ala Thr Thr Asn Thr Trp Ser Val Ser Lys Glu Ser Phe Ile  
Asp Tyr Leu Ser Lys Val Ile Ile Thr Val Lys Asp Val Asn Asp Gln Ile Val Phe Thr Lys Glu Thr  
Thr Asn Lys Thr Asn Ile Tyr Phe Glu Ile Glu Leu Leu Pro Gly Thr Tyr Thr Phe Glu Val Lys Gly  
Tyr Glu Glu Asp Leu Val Ile Phe Ser Gly Glu Lys Val Asn Gln Ile Ile Asp Glu Lys Asn Asn Ile  
Val Asn Val Glu Thr Phe Phe Val Asn Gly Ile Val Arg Thr Ile Ile Glu Val Asp Asp Ile Ile Tyr Lys  
Asn Tyr Asp Ile Thr Ser Ala Thr Leu Ile Phe Lys Lys Asp Thr Ala Gln Glu Asp Tyr Glu Glu Val  
Pro Val Thr Leu Thr Gly Thr Ser Thr Leu Ile Asn Lys Glu Leu Tyr Pro Gly Met Trp Thr Val Lys  
Phe Glu Val Asp Leu Lys Ser Lys Asp Ala Ser Met Leu Pro Glu Lys Val His Leu Glu Asn Glu Phe  
Ser Ile Glu Val Leu Pro Ala Lys Thr Lys Ser Leu Thr Phe Asn Val Val Phe Asp Thr Glu Val Asn  
Glu Pro Lys Leu Val Val Val Phe Pro Gln Ile Glu Leu Pro Phe Val Asp Pro Val Thr Asn Leu Ser  
Gly Glu Ile Asn Glu Leu Glu Gly Asn Leu Ser Met Asn Trp Asp Tyr Ser Asp Pro Asn Ala Glu Phe  
Tyr Val Tyr Lys Glu Leu Glu Glu Gln Gly Glu Tyr Leu Tyr Glu Phe Val Gly Lys Thr Arg Glu Lys  
Ser Tyr Thr Ile Glu Asn Phe Thr Lys Gln Glu Phe Asp Lys Phe Ser Gly Ile Ala Ile Asn Val Tyr  
Ala Asn Gly Lys Glu Ser Gly Leu Val Val Leu Lys Lys Glu Asn Ile Lys Leu Ile Asp Leu Glu Ser  
Val Asp Ser Ile Ser Ala Thr Tyr Asn Val Asp Thr Asn Glu Leu Lys Leu Asp Trp Asn Tyr Thr Asn  
Ser Ser Val Thr Phe Glu Val Leu Lys Lys Gly Ile Asn Ser Asn Glu Tyr Glu Ile Ile Ser Gln Leu Thr  
Gln Asn Ser Phe Ser Thr Glu Phe Thr Gly Arg Gln Phe Trp Asp Leu Glu Lys Ile Ala Ile Arg Val  
Val Ala Asn Gly Phe Glu Ser Lys Ile Asn Glu Ile Ser Arg Asp Asp Ile Thr Ile Thr Ser Leu Asn Leu  
Pro Leu Thr Ser Ser Thr Met Tyr Thr Leu Phe Ile Arg Ser Tyr Phe Asp Thr Asp Gly Asp Gly Val  
Gly Asp Phe Ser Gly Val Ala Glu Lys Val Asp Tyr Leu Lys Ser Leu Gly Val Asp Thr Val Trp Phe  
Leu Pro Phe Asn Lys Ser Lys Ser Tyr His Gly Tyr Asp Val Glu Asp Tyr Tyr Asp Val Glu Pro Asp  
Tyr Gly Thr Leu Gln Asp Leu Asp Asn Met Ile Lys Val Leu Asn Glu Asn Gly Ile Lys Val Val Met  
Asp Leu Val Val Asn His Thr Ser Asp Thr His Pro Trp Phe Leu Asp Ala Val Glu Asn Thr Thr Asn  
Ser Pro Tyr Trp Asn Tyr Tyr Ile Met Ser Leu Asp Glu Pro Gln Asn Lys Asn His Trp His Tyr Lys  
Val Asn Ser Lys Gly Gln Thr Val Trp Tyr Phe Gly Leu Phe Asp Ser Ser Met Pro Asp Leu Asn Tyr  
Asp Asn Pro Lys Val Met Asp Glu Val Lys Lys Ile Ile Asp Phe Trp Ala Asp Met Gly Val Asp Gly  
Phe Arg Leu Asp Ala Ala Lys His Tyr Tyr Gly Phe Asp Trp Ser Asp Gly Ile Glu Gln Ser Ala Ser  
Val Ala Lys Glu Ile Glu Asp Tyr Ile Lys Asp Lys Leu Gly Glu Asn Ala Ile Val Val Ser Glu Val

Figure 16 (cont.)

Tyr Asp Gly Asp Ser Asn Val Leu Leu Lys Phe Ala Pro Met Pro Val Phe Asn Phe Ser Phe Met Tyr  
Asn Leu Arg Gly Asn Phe Glu Gly Arg Asp Asn Leu Ile Ser Asp Ser Ile Ser Trp Val Asp Ser Ser  
Leu Tyr Asn Leu Asn Val Phe His Phe Pro Phe Ile Asp Ser His Asp Leu Asp Arg Phe Ile Ser Glu  
Leu Val Asp Ser Lys Tyr Gln Gly Asp Val Ile Ser Ala Thr Lys Gln Tyr Leu Leu Val Asn Ala Leu  
Leu Leu Ser Leu Thr Gly Met Pro Thr Ile Tyr Tyr Gly Asp Glu Ile Gly Leu Arg Gly Trp Lys Trp  
His Ser Glu Pro Trp Asp Ile Pro Val Arg Glu Pro Met Gln Trp Tyr Lys Asp Gln Lys Gly Asn Gly  
Gln Thr Tyr Trp Thr Lys Glu Phe Tyr Glu Gly Ile Thr Glu Gly Ser Ala Asn Glu Asp Gly Ala Ile  
Tyr Asp Asp Pro Asp Asp Gly Val Ser Val Glu Glu Gln Glu Asn Gly Tyr Ser Ile Leu Asn Phe Phe  
Lys Glu Phe Ile Asn Leu Arg Lys Asp Tyr Pro Ala Leu Ala Phe Gly Ser Thr Thr Ile Glu Arg Asp  
Trp Lys Asn Leu Tyr Val Leu Lys Lys Ser Tyr Asn Phe Gln Asp Val Leu Val Leu Ile Asn Leu Asp  
Pro Thr Tyr Ser Asn Thr Tyr Glu Val Pro Glu Gly Tyr Lys Trp Val Trp Tyr Ala Phe Phe Asp Gly  
Asp Asn Tyr Glu Phe Gly Ala Lys Asp Glu Met Ile Leu Gln Asn Thr Ser Trp Thr Ile Asn Pro Arg  
Gln Ile Tyr Ile Phe Val Lys

SEQ ID NO: 99

atgtacacactcttcatccgctctttttacgatacaacaacgacggtgtaggtgactacaacggtgtgccccaaaaagtagactatctcaaacg  
cttgagtggtacacagtttggttcttgcggttcaacaaagcaaaatcgaccacggttacgatgtgaagactactacgatgtgaacctgactatg  
gaacatacgcaacaacttgaataatgataaagacactcaatcagaacggaattcggttggacttgggttggaaccacacttccgatacac  
actcgtggttcttgatgccgttgagaacacaacgaattcgaaatttgagctactacataatgacacttgaataagagacggttgaatcact  
ggcattggaagataaactcaaaagggcaaaagtttactacttcggactgttgactcatcaatgcccgatttgaatttcgacaatccacaagtgat  
gaacgaaatcaagagaataatcgatttctggataacagttggtgtggtggttcagacttgatgcacaaagcactacaaaggctgggattggg  
acgacggcatttcaggttcagcagcaatcgcgagggaatagaaagtacatcaggagcaagttaggaacgatgcgatagttgcggggaa  
gtgtacgatggaatccatcggttctttcacaatttcacccgatgccggcggttcaacttcacattcatgtatggaataacaggcaacctgagggg  
aaagataacctgctgggagaaacaatttcattggttaattggagcgagttattatctcaacgtaaaacatttcccgttcatagacaatcacgattga  
acagatggatcgcatacttatcgacaaaagtatatggaacacacaagttggtacgaagcagtatatttaacaaatgcgctcttgccttctta  
aacggtatgcctgttatttattatgggaatgaaataggcttgagaggtggaatggggacaagaccggtgggatttgcgggtgagagagccga  
tgcagtggtacgcaagtcaaaagtgagctgggcagacatggtggacaagcctgcttaccagcaaaaggaatcacatttggaatgcaaac  
gtcagtggtgcgatgtacgatgatccaatgatggggttcagtagaagagcagatgaatggttacacgataaataacttcttaaacaaatcataa  
ccctgaggaagacatatccggctctatcgaaagggttcgataacgatagaacgcgactggaagaacctgtacgttatcaaacgagctacggaa  
atcaggaagtgcctgtattgataaacttagaccaacttgccgaacaattacacgttaccaggtggatacaggtgggtctggtatgcgttcttaa  
tgggagttgttgaatttggcaataaaaacgaatcaccactgagccaagataccaactggacagtcattcaaggcaagtgatgtgttgtgaa  
ggactaa

SEQ ID NO: 100

Met Tyr Thr Leu Phe Ile Arg Ser Phe Tyr Asp Thr Asn Asn Asp Gly Val Gly Asp Tyr Asn Gly Val  
Ala Gln Lys Val Asp Tyr Leu Lys Thr Leu Gly Val Asp Thr Val Trp Phe Leu Pro Phe Asn Lys Ala  
Lys Ser Tyr His Gly Tyr Asp Val Glu Asp Tyr Tyr Asp Val Glu Pro Asp Tyr Gly Thr Tyr Ala Gln  
Leu Glu Asn Met Ile Lys Thr Leu Asn Gln Asn Gly Ile Arg Val Val Met Asp Leu Val Val Asn His  
Thr Ser Asp Thr His Ser Trp Phe Leu Asp Ala Val Glu Asn Thr Thr Asn Ser Lys Tyr Trp Ser Tyr  
Tyr Ile Met Thr Leu Glu Asn Arg Asp Gly Trp Asn His Trp His Trp Lys Ile Asn Ser Lys Gly Gln  
Lys Val Tyr Tyr Phe Gly Leu Phe Asp Ser Ser Met Pro Asp Leu Asn Phe Asp Asn Pro Gln Val  
Met Asn Glu Ile Lys Arg Ile Ile Asp Phe Trp Ile Thr Val Gly Val Asp Gly Phe Arg Leu Asp Ala  
Pro Lys His Tyr Lys Gly Trp Asp Trp Asp Asp Gly Ile Ser Gly Ser Ala Ala Ile Ala Arg Glu Ile Glu  
Ser Tyr Ile Arg Ser Lys Leu Gly Asn Asp Ala Ile Val Val Gly Glu Val Tyr Asp Gly Asn Pro Ser  
Val Leu Ser Gln Phe Ala Pro Met Pro Ala Phe Asn Phe Thr Phe Met Tyr Gly Ile Thr Gly Asn His  
Glu Gly Lys Asp Asn Leu Leu Gly Glu Thr Ile Ser Trp Val Asn Gly Ala Ser Tyr Tyr Leu Asn Val  
Lys His Phe Pro Phe Ile Asp Asn His Asp Leu Asn Arg Trp Ile Ser Ile Leu Ile Asp Gln Lys Tyr Ser  
Gly Asn Thr Gln Val Gly Thr Lys Gln Tyr Ile Leu Thr Asn Ala Leu Leu Leu Ser Leu Asn Gly Met  
Pro Val Ile Tyr Tyr Gly Asn Glu Ile Gly Leu Arg Gly Trp Lys Trp Gly Gln Asp Pro Trp Asp Leu  
Pro Val Arg Glu Pro Met Gln Trp Tyr Ala Ser Gln Ser Gly Ala Gly Gln Thr Trp Trp Thr Lys Pro

Figure 16 (cont.)

Val Tyr Gln Gln Lys Gly Ile Thr Phe Gly Asn Ala Asn Val Asp Gly Ala Met Tyr Asp Asp Pro Asn  
Asp Gly Val Ser Val Glu Glu Gln Met Asn Gly Tyr Thr Ile Asn Asn Phe Phe Lys Gln Phe Ile Thr  
Leu Arg Lys Thr Tyr Pro Ala Leu Ser Lys Gly Ser Ile Thr Ile Glu Arg Asp Trp Lys Asn Leu Tyr  
Val Ile Lys Arg Val Tyr Gly Asn Gln Glu Val Leu Val Leu Ile Asn Leu Asp Pro Thr Trp Pro Asn  
Asn Tyr Thr Leu Pro Gly Gly Tyr Arg Trp Val Trp Tyr Ala Phe Phe Asn Gly Ser Leu Phe Glu Phe  
Gly Asn Lys Asn Glu Ser Pro Leu Ser Gln Asp Thr Asn Trp Thr Val Asn Pro Arg Gln Val Tyr Val  
Phe Val Lys Asp

SEQ ID NO: 101

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aaccgctttgtacacctctttgaatggaaatgggaagatgttcacaggagtgtaaacatttctcggacctaaggtttgccgcagtgcaagt  
ctctccgccaactaaatctcaaacacggatgcatgggtggggccgttatcaaccggttagttatgctttgaaggacgcagcggtaatcgagcc  
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ctgatgtaccctatagcagtaatgactttaactcctgtacaggagatattgactataataaccgttggcaaacacagcattgtgattagtcggtctta  
atgatctaaaaacaggatctgactacgtccgcaaaaaatagcggattatgaacgacgcaatcagtatgggtgtagctggtttccgtattgatg  
cagccaaacatataccagcaggtgatatagctgccattaaaggtaaattaaatggtaatccatacatctccaagaggtaattgggtgcatccggcg  
aacctgttcgaccgactgaatacacctttatcggtgtgcacggaatttcaatttgcgcgaaattgggtccagcctccgcaatagtaattgctt  
gggtaaaagacattggcagtcacaaatgaattatccagtgctgatgccgaacatttgaacgaatcatgatgaagagcgtcataaccggaatggtc  
ctatttggcagcggcgttcaaggtaattgggtatgcattagcaaatattttcaccttagcttaccctacggctatccaaaaatcatgtcaggatacttctt  
ccacgggtgactttaacgcagctccaccaagcagtggtatcacacaggaaatgcgtgtggtttgatggcggagactgggtatgcgaacacaa  
atggcgcggtattgctaacatgggtgccttccgcaactatacagcaagcgaatggcgtatcagtaattgggtggcaaacagtaacgaccaaattg  
cttttggcgcggtgggttaggtttgtgttattaataaacgtgctaattggttagcattaatacaagttttgatacgggaatgcctgatggccaactgt  
aacataatagaagtaactttgatgaaagcaccggccaatgtagtgcagctacagattccaacgggtcaagcgttattaccgtcagtggtgggca  
agctaactttaatgtagcaggcgatcatgctgtgcattcatgttggcgcaaaaattgggtgatcaatgtatggtgatgattgccatgtacagga  
tccgattgtaataatgatctaaacctgatttgcagtaccagcaacatcaattgtacatcagaaaattacctaactgctatattactggggagcaca  
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aaattaacgccatcttagtgacaatgggtgcaataaaacagctgatctaactgttactgggtgcaggtgttataaagacgggacttggagcacctt  
acaaaattgtggcttgaattaccgggtgcacaaaccaatccagtcgggtggcgacgaagtctggtacttccgaggtactgtaataactgactggggtta  
aagcacaattagattatgacgcaactagcgggttattacacatacaaaagctttaaattggtgaagaagcacctgcgcgtttaaattgataatgggt  
agttggactgaagcttatccaacagctgattaccaagttacagataacaattcataccgcattaacttaataagcgatagcaaacgattacagtaa  
acgcacaataa

SEQ ID NO: 102

Met Arg Phe Phe Pro Lys Leu Ile Ser Pro Phe Pro Gln Asn Thr Arg Glu Trp Gln Arg Ser Ala Val  
Ser Arg Asp Thr Glu Gln Leu Gln Arg Lys Val Ile Met Ile Asn Leu Lys Lys Asn Thr Ile Ser Ala  
Leu Val Ala Gly Met Val Leu Gly Phe Ala Ser Asn Ala Met Ala Val Pro Arg Thr Ala Phe Val His  
Leu Phe Glu Trp Lys Trp Glu Asp Val Ala Gln Glu Cys Glu Thr Phe Leu Gly Pro Lys Gly Phe Ala  
Ala Val Gln Val Ser Pro Pro Thr Lys Ser His Asn Thr Asp Ala Trp Trp Gly Arg Tyr Gln Pro Val  
Ser Tyr Ala Phe Glu Gly Arg Ser Gly Asn Arg Ser Gln Phe Lys Asn Met Val Gln Arg Cys Lys Ala  
Val Gly Val Asp Ile Tyr Val Asp Ala Val Ile Asn His Met Ala Ala Tyr Asp Arg Asn Phe Pro Asp  
Val Pro Tyr Ser Ser Asn Asp Phe Asn Ser Cys Thr Gly Asp Ile Asp Tyr Asn Asn Arg Trp Gln Thr  
Gln His Cys Asp Leu Val Gly Leu Asn Asp Leu Lys Thr Gly Ser Asp Tyr Val Arg Gln Lys Ile Ala  
Asp Tyr Met Asn Asp Ala Ile Ser Met Gly Val Ala Gly Phe Arg Ile Asp Ala Ala Lys His Ile Pro  
Ala Gly Asp Ile Ala Ala Ile Lys Gly Lys Leu Asn Gly Asn Pro Tyr Ile Phe Gln Glu Val Ile Gly Ala  
Ser Gly Glu Pro Val Arg Pro Thr Glu Tyr Thr Phe Ile Gly Gly Val Thr Glu Phe Gln Phe Ala Arg  
Lys Leu Gly Pro Ala Phe Arg Asn Ser Asn Ile Ala Trp Leu Lys Asp Ile Gly Ser Gln Met Glu Leu  
Ser Ser Ala Asp Ala Val Thr Phe Val Thr Asn His Asp Glu Glu Arg His Asn Pro Asn Gly Pro Ile  
Trp His Gly Val Gln Gly Asn Gly Tyr Ala Leu Ala Asn Ile Phe Thr Leu Ala Tyr Pro Tyr Gly Tyr  
Pro Lys Ile Met Ser Gly Tyr Phe Phe His Gly Asp Phe Asn Ala Ala Pro Pro Ser Ser Gly Ile His Thr

Figure 16 (cont.)

Gly Asn Ala Cys Gly Phe Asp Gly Gly Asp Trp Val Cys Glu His Lys Trp Arg Gly Ile Ala Asn Met  
Val Ala Phe Arg Asn Tyr Thr Ala Ser Glu Trp Arg Ile Ser Asn Trp Trp Gln Asn Ser Asn Asp Gln  
Ile Ala Phe Gly Arg Gly Gly Leu Gly Phe Val Val Ile Asn Lys Arg Ala Asn Gly Ser Ile Asn Gln  
Ser Phe Asp Thr Gly Met Pro Asp Gly Gln Tyr Cys Asn Ile Ile Glu Ala Asn Phe Asp Glu Ser Thr  
Gly Gln Cys Ser Ala Ala Thr Asp Ser Asn Gly Gln Ala Val Ile Thr Val Ser Gly Gly Gln Ala Asn  
Phe Asn Val Ala Gly Asp His Ala Ala Ala Ile His Val Gly Ala Lys Ile Gly Asp Gln Cys Ser Gly  
Asp Asp Cys Pro Cys Thr Gly Ser Asp Cys Asn Asn Asp Pro Lys Pro Asp Phe Ala Val Pro Ala  
Thr Ser Ile Cys Thr Ser Glu Asn Leu Pro Thr Leu Tyr Tyr Trp Gly Ala Gln Pro Thr Asp Ser Leu  
Ala Asn Ala Ala Trp Pro Gly Val Ala Met Gln Thr Asn Gly Asp Phe Lys Cys His Asp Leu Gly Val  
Glu Leu Thr Lys Ile Asn Ala Ile Phe Ser Asp Asn Gly Ala Asn Lys Thr Ala Asp Leu Thr Val Thr  
Gly Ala Gly Cys Tyr Lys Asp Gly Thr Trp Ser Thr Leu Gln Asn Cys Gly Phe Glu Ile Thr Gly Ala  
Gln Thr Asn Pro Val Gly Gly Asp Glu Val Trp Tyr Phe Arg Gly Thr Ala Asn Asp Trp Gly Lys Ala  
Gln Leu Asp Tyr Asp Ala Thr Ser Gly Leu Tyr Tyr Thr Ile Gln Ser Phe Asn Gly Glu Glu Ala Pro  
Ala Arg Phe Lys Ile Asp Asn Gly Ser Trp Thr Glu Ala Tyr Pro Thr Ala Asp Tyr Gln Val Thr Asp  
Asn Asn Ser Tyr Arg Ile Asn Phe Asn Ser Asp Ser Lys Ala Ile Thr Val Asn Ala Gln

SEQ ID NO: 103

gtgctaacgtttcaccgcatcattcgaaaaggatggatgttctgctcgctgttttgcactgcctcgctgttctgcccacaggacagcccgcca  
aggctgccgcaccgtttaacggcaccatgatgcagtattttgaatggtacttgccggatgatggcacgttatggaccaaagtggccaatgaagc  
caacaactatccagccttgccatcaccgctcttttggtgctgccgccgcttacaaaggaacaagccgcagcgacgtagggtacggagtatacga  
ctgtatgacctcgccgaattcaatcaaaaaggacgctccgcacaaaatacggacaaaagctcaatatctcaagccattcaagccgcccac  
gccgctggaatgcaagtgtacgccgatgtcgtgttcgaccataaaggcgccgacggcacggaatgggtggacgccgtcgaagtcaatc  
cgtccgaccgcaaccaagaatctcgggcacctatcaaatccaagcatggacgaaatttgatttcccgggcgggggcaacacctactccagctt  
taagtggcgctggttaccattttgacggcgttgattgggacgaaagccgaaaattgagccgattacaaattccgcggcatcgccaaagcgtgg  
gattgggaagtagacacggaaaacgaaactatgactacttaatgtatgccgaccttgataggatcatcccgaagtcgtgaccgagctgaaaa  
actgggggggaatggtatgtcaacacaacgaacattgatgggtccggcttgatgccgtcaagcataattaagttcagtttttctgattggtgtcgt  
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gaacgatgtctttgtttgatgccccgttacacaacaaattttataccgctccaaatcaggggcgcatattgatatgcgcaggttaataccaataact  
ctcatgaaagatcaaccgacattggccgctaccttcgttgataatcatgacaccgaacccggccaagcgtgcagtcagtggtcgacccatggt  
tcaaaccgttggttacgcctttattctaactcggcaggaagatacccgctgcgtcttttatggtgactattatggcattccacaataataacattccttc  
gctgaaaagcaaaatcgatccgctcctcatcgcgcgcagggtattatgcttacggaacgcaacatgattatcttgatcactccgacatcatcggt  
ggacaagggaagggtcactgaaaaccaggatccgggctggccgactgatcccgatgggcccgggaggaagcaaatggatgtacgttg  
gcaacaacacgctggaaaagtgtctatgacctaccggcaaccggagtacaccgtcaccatcaacagtgatggatggggggaattcaaa  
gtcaatggcggttcggttcggttggttcctagaaaaacgacctgttctaccatcgctcgccgatcacaacccgacctggactggtgaattc  
gtccgttgaccgaaccacggttggtggcatggccttga

SEQ ID NO: 104

Val Leu Thr Phe His Arg Ile Ile Arg Lys Gly Trp Met Phe Leu Leu Ala Phe Leu Leu Thr Ala Ser  
Leu Phe Cys Pro Thr Gly Gln Pro Ala Lys Ala Ala Ala Pro Phe Asn Gly Thr Met Met Gln Tyr Phe  
Glu Trp Tyr Leu Pro Asp Asp Gly Thr Leu Trp Thr Lys Val Ala Asn Glu Ala Asn Asn Leu Ser Ser  
Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys Gly Thr Ser Arg Ser Asp Val Gly Tyr Gly  
Val Tyr Asp Leu Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr  
Lys Ala Gln Tyr Leu Gln Ala Ile Gln Ala Ala His Ala Ala Gly Met Gln Val Tyr Ala Asp Val Val  
Phe Asp His Lys Gly Gly Ala Asp Gly Thr Glu Trp Val Asp Ala Val Glu Val Asn Pro Ser Asp Arg  
Asn Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn  
Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr His Phe Asp Gly Val Asp Trp Asp Glu Ser Arg Lys Leu  
Ser Arg Ile Tyr Lys Phe Arg Gly Ile Gly Lys Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn  
Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp His Pro Glu Val Val Thr Glu Leu Lys Asn  
Trp Gly Glu Trp Tyr Val Asn Thr Thr Asn Ile Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys  
Phe Ser Phe Phe Pro Asp Trp Leu Ser Tyr Val Arg Ser Gln Thr Gly Lys Pro Leu Phe Thr Val Gly

Figure 16 (cont.)

Glu Tyr Trp Ser Tyr Asp Ile Asn Lys Leu His Asn Tyr Ile Thr Lys Thr Asn Gly Thr Met Ser Leu  
Phe Asp Ala Pro Leu His Asn Lys Phe Tyr Thr Ala Ser Lys Ser Gly Gly Ala Phe Asp Met Arg Thr  
Leu Met Thr Asn Thr Leu Met Lys Asp Gln Pro Thr Leu Ala Val Thr Phe Val Asp Asn His Asp  
Thr Glu Pro Gly Gln Ala Leu Gln Ser Trp Val Asp Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile  
Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly Asp Tyr Tyr Gly Ile Pro Gln Tyr Asn Ile  
Pro Ser Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln His  
Asp Tyr Leu Asp His Ser Asp Ile Ile Gly Trp Thr Arg Glu Gly Val Thr Glu Lys Pro Gly Ser Gly  
Leu Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln His Ala Gly  
Lys Val Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn Ser Asp Gly Trp Gly Glu  
Phe Lys Val Asn Gly Gly Ser Val Ser Val Trp Val Pro Arg Lys Thr Thr Val Ser Thr Ile Ala Arg  
Pro Ile Thr Thr Arg Pro Trp Thr Gly Glu Phe Val Arg Trp Thr Glu Pro Arg Leu Val Ala Trp Pro

SEQ ID NO: 105

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ttagcgaatcttggcattactgcccttggcttccccctgcctataaaggaacaagcagcagtgacgttgatattggcgtttatgatttatgacct  
aggagagttaatacaaaaaggaactgtccgaacaaaatacgaacaaaaacacataatccaagcaatccaagcggcgcatacagcaggaa  
tgcaagtatatgcagatgctgctttaaccataaagccggtgcagatgggacagaactagtggatgcagtagaagtaaaccctctgaccgcaat  
caagaaatatcaggaacatatacaatccaagcgtggacaaaattgatttctggtcgtggaacaccattctagttttaaatggcgttggtatca  
ttcgtatggaacggactgggatgagagtagaaaaactaaatcgtattacaattccgcggcacgggaaaagcatgggattgggaagtagataca  
gaaaatgggaattatgactatctcatgtatgcagatttggatattgatcatccagagggtgtatctgaactaaaaattggggaaaagtggtatgtaa  
ccacaaccaatcgcaggtatccgctggtgcagtgagcatattaaatatacttttccagactggctatcgtatgtacgaacccaaacac  
aaaagcctcttttggcgttggcggaatttggagctatgcattaacaagctacacaactataatacaagacgaacggctctatgtccctattcgat  
gccccgctgcataacaattttatagcatcgaaatcaggtggctatttggatgctgcacattactcaacaacacattgatgaaagtaacca  
cactatcggtcacattagtagacaatcacgatactgagccagggaatctttgcagtcgtgggtcagccgtggtttaaaccgttagcttacgcat  
ttatctgaccgcgaagaagggtatccgtgcacatctttatggagattactatggtattccaaaatacaacattcctgcgctgaaaagcaaacctgac  
cgctgttaattgctcgaagagattatgcctacggaacacagcagcactatattgacaatgcagatatcggctggacgcgggaaggagtagct  
gaaaaagcaaatcgggacttgcgcactcattaccgacggacctggcggaagcaaatggatgtatgttgcaaacaacacgctggcaaac  
gtttatgatctaaccggcaatcgaagtatacagtgacaatcaacgctgatggatggggagaatttaaagtcaatggagggtctgtatccatag  
ggttccaaaacatcaaccacttcccaatcacatttactgtaataatgccacaaccgttggggacaaaatgtatacgttgctcggaatatttcg  
cagctgggcaac

SEQ ID NO: 106

Met Ser Leu Phe Lys Lys Ile Phe Pro Trp Ile Val Ser Leu Leu Leu Leu Phe Ser Phe Ile Ala Pro Phe  
Ser Ile Gln Thr Glu Lys Val Arg Ala Gly Ser Val Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu  
Trp Tyr Leu Pro Asp Asp Gly Thr Leu Trp Thr Lys Val Ala Asn Asn Ala Gln Ser Leu Ala Asn Leu  
Gly Ile Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys Gly Thr Ser Ser Ser Asp Val Gly Tyr Gly Val  
Tyr Asp Leu Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr Lys  
Thr Gln Tyr Ile Gln Ala Ile Gln Ala Ala His Thr Ala Gly Met Gln Val Tyr Ala Asp Val Val Phe  
Asn His Lys Ala Gly Ala Asp Gly Thr Glu Leu Val Asp Ala Val Glu Val Asn Pro Ser Asp Arg Asn  
Gln Glu Ile Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn Thr  
Tyr Ser Ser Phe Lys Trp Arg Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn  
Arg Ile Tyr Lys Phe Arg Gly Thr Gly Lys Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn Tyr  
Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp His Pro Glu Val Val Ser Glu Leu Lys Asn Trp  
Gly Lys Trp Tyr Val Thr Thr Thr Asn Ile Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Tyr  
Ser Phe Phe Pro Asp Trp Leu Ser Tyr Val Arg Thr Gln Thr Gln Lys Pro Leu Phe Ala Val Gly Glu  
Phe Trp Ser Tyr Asp Ile Asn Lys Leu His Asn Tyr Ile Thr Lys Thr Asn Gly Ser Met Ser Leu Phe  
Asp Ala Pro Leu His Asn Asn Phe Tyr Ile Ala Ser Lys Ser Gly Gly Tyr Phe Asp Met Arg Thr Leu  
Leu Asn Asn Thr Leu Met Lys Asp Gln Pro Thr Leu Ser Val Thr Leu Val Asp Asn His Asp Thr

Figure 16 (cont.)

Glu Pro Gly Gln Ser Leu Gln Ser Trp Val Glu Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu  
Thr Arg Gln Glu Gly Tyr Pro Cys Ile Phe Tyr Gly Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro  
Ala Leu Lys Ser Lys Leu Asp Pro Leu Leu Ile Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln His Asp  
Tyr Ile Asp Asn Ala Asp Ile Ile Gly Trp Thr Arg Glu Gly Val Ala Glu Lys Ala Asn Ser Gly Leu  
Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln His Ala Gly Lys  
Thr Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn Ala Asp Gly Trp Gly Glu Phe  
Lys Val Asn Gly Gly Ser Val Ser Ile Trp Val Pro Lys Thr Ser Thr Thr Ser Gln Ile Thr Phe Thr Val  
Asn Asn Ala Thr Thr Val Trp Gly Gln Asn Val Tyr Val Val Gly Asn Ile Ser Gln Leu Gly Asn

SEQ ID NO: 107

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cgaccgtttccaggatggcgacaagaccaacaacatggacgtgggtcccgacggacatgaaaaatatcatggcggcgacatccaggggctc  
atcgacaagctcgactatatcaaggagaccgggttcgacggccatctggctcacgccccctatgaaggggcagacccacttcttcgagaccgac  
aattaccatggttactggccccattgacttctatgacacggacccccatgtgggcaccatgcagaaatttgaggagcttatcgagaaagcccatga  
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accatataggagatgtgaaggactgggaagatccctactgggctgaaaacggctccatattcgggtcttctgacctggcgagggaaaacctg  
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cgggagggcagaccccgaaggacgagctctatcgtggtgcttaacaacggctatgatacgaggaacgggacataccgctccgccccgagag  
cggcatcaagaacggcacggtgctgaaggatgcatcaccggcgaaccgtgacggtacagaacggaaaaatccatgcgaaatgcggcg  
caaacaggcgcggtatctacgtgccccgctag

SEQ ID NO: 108

Met Asp Ser Leu Asp Ala Pro Glu Gln Lys Pro Trp Val Lys Asp Gly Arg Leu Ser Ala Tyr Leu Asp  
Thr Gly Thr Gly Thr Val Val Ala Pro Glu Ala Pro Ala Pro Pro Pro Ala Glu Glu Val Arg  
Pro Val Asp Lys Trp Lys Asn Asp Ile Ile Tyr Phe Val Leu Thr Asp Arg Phe Gln Asp Gly Asp Lys  
Thr Asn Asn Met Asp Val Val Pro Thr Asp Met Lys Lys Tyr His Gly Gly Asp Ile Gln Gly Leu Ile  
Asp Lys Leu Asp Tyr Ile Lys Glu Thr Gly Ser Thr Ala Ile Trp Leu Thr Pro Pro Met Lys Gly Gln  
Thr His Phe Phe Glu Thr Asp Asn Tyr His Gly Tyr Trp Pro Ile Asp Phe Tyr Asp Thr Asp Pro His  
Val Gly Thr Met Gln Lys Phe Glu Glu Leu Ile Glu Lys Ala His Glu Lys Gly Leu Lys Ile Val Leu  
Asp Ile Pro Leu Asn His Thr Ala Trp Glu His Pro Phe Tyr Lys Asp Asp Ser Lys Lys Asp Trp Phe  
His His Ile Gly Asp Val Lys Asp Trp Glu Asp Pro Tyr Trp Ala Glu Asn Gly Ser Ile Phe Gly Leu  
Pro Asp Leu Ala Gln Glu Asn Pro Ala Val Glu Lys Tyr Leu Ile Asp Val Ala Lys Phe Trp Val Asp  
Lys Gly Ile Asp Gly Phe Arg Leu Asp Ala Val Lys Asn Val Pro Leu Asn Phe Trp Ala Lys Phe Asp  
Arg Ala Ile His Asp Tyr Ala Gly Lys Asp Phe Leu Leu Val Gly Glu Tyr Phe Asp Gly Asn Pro Ala  
Lys Val Ala Asn Tyr Gln Arg Glu Asp Met Ser Ser Leu Phe Asp Tyr Pro Leu Tyr Trp Thr Leu Lys  
Asp Thr Phe Ala Lys Asp Gly Ser Met Arg Asn Leu Ala Ala Lys Leu Asp Glu Cys Asp Arg Asn  
Tyr Pro Asp Pro Gly Leu Met Ser Val Phe Leu Asp Asn His Asp Thr Pro Arg Phe Leu Thr Glu Ala  
Asn Gly Asn Lys Asp Lys Leu Lys Leu Ala Leu Ala Phe Ala Met Thr Ile Asn Arg Met Pro Thr Ile  
Tyr Tyr Gly Thr Glu Val Ala Met Glu Gly Asn Cys Asp Ile Met Gly Ala Val Asp Asn Arg Arg  
Asp Met Gln Trp Asp Lys Asp Pro Asp Met Phe Lys Tyr Phe Lys Thr Leu Thr Thr Ala Arg Asn  
Glu His Glu Ser Leu Arg Glu Gly Lys Lys Leu Glu Met Trp Gln Asp Asp Lys Val Tyr Ala Tyr Gly  
Arg Gln Thr Pro Lys Asp Glu Ser Ile Val Val Leu Asn Asn Gly Tyr Asp Thr Gln Glu Arg Asp Ile

Figure 16 (cont.)

Pro Leu Arg Pro Glu Ser Gly Ile Lys Asn Gly Thr Val Leu Lys Asp Val Ile Thr Gly Glu Thr Val  
Thr Val Gln Asn Gly Lys Ile His Ala Lys Cys Gly Gly Lys Gln Ala Arg Ile Tyr Val Pro Ala

SEQ ID NO: 109

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ctcagcgatatggattccaccagcgagtaagggcatgagcgggtgttattccatgggctacgatccctacgatttcttgacctcggcgagtacta  
tcagaagggggacagttgagacgcgcttcggctcaaggaagaactggtgaacatgataaacaccgcacactcctacggcataaaggatag  
cggacatagtcataaaccaccgcgcgggtggagaccttgagtgaacccctcgtgaacgactataacctggacagacttctcaaaagtcgcctc  
cggtaaatatagggccaactacctgacttccacccaaacgagcttactgtgtgatgaaggtaccttggaggataacctgatatatgtcacga  
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gcatccacgagtagaccggcaacctcggcggttgatagacaagtacgtctcctccagcgggtgggtctatctgaggccccagcccacgac  
ccggcgaacggctactacggctactctgtctggagctactgcggtgtgggtga

SEQ ID NO: 110

Met Ala Arg Lys Thr Leu Ala Ile Phe Phe Val Leu Leu Val Leu Leu Ser Leu Ser Ala Val Pro Ala  
Lys Ala Glu Thr Leu Glu Asn Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Gly Gly Gly  
Ile Trp Trp Asp Thr Ile Ala Gln Lys Ile Pro Glu Trp Ala Ser Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro  
Ala Ser Lys Gly Met Ser Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp Leu Gly Glu  
Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Glu Glu Leu Val Asn Met Ile Asn Thr  
Ala His Ser Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp Leu Glu Trp  
Asn Pro Phe Val Asn Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn  
Tyr Leu Asp Phe His Pro Asn Glu Leu His Cys Cys Asp Glu Gly Thr Phe Gly Gly Tyr Pro Asp Ile  
Cys His Asp Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Ser Glu Ser Tyr Ala Ala Tyr Leu Arg  
Ser Ile Gly Val Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val Asn Asp Trp  
Leu Ser Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Leu Leu Asn Trp  
Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp Asn  
Thr Asn Ile Pro Ala Leu Val Asp Ala Leu Arg Tyr Gly Gln Thr Val Val Ser Arg Asp Pro Phe Lys  
Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile  
Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Lys Leu  
Asn Asn Leu Ile Trp Ile His Asp His Leu Ala Gly Gly Ser Thr Asp Ile Val Tyr Tyr Asp Ser Asp  
Glu Leu Ile Phe Val Arg Asn Gly Tyr Gly Thr Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Gly Ser  
Ser Lys Val Gly Arg Trp Val Tyr Val Pro Lys Phe Ala Gly Ser Cys Ile His Glu Tyr Thr Gly Asn  
Leu Gly Gly Trp Ile Asp Lys Tyr Val Ser Ser Ser Gly Trp Val Tyr Leu Glu Ala Pro Ala His Asp  
Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Cys Gly Val Gly

SEQ ID NO: 111

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cataagacagaaaatcccgagtggtacgacgctggaatctcggcgatatgattctccagctagcaaaggtatgggtggtgcatactccatg  
ggttatgaccttaccgatttcttgacctcggcgagtactatcagaagggaacagttgagacgcgcttcggctcaaggagggaactggtgaaca  
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taaacaactatacttgacagacttctcaaggctcgcctccggtaaatacagggccaactaccttacttccacccaaacgaggtcaagtgctgc  
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Figure 16 (cont.)

tatctccggagcatagggatcgatgcatggcggttcgactacgtcaaagggttacggagcgtgggttgtaacgactggctcagctgggtggggag  
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ctactacgataacgatgagctaataatcatgagggagggctacgggagcaagccgggctcataacctacataaaacctcgaaacgactggg  
ccgagcgtgggtgaacgtcggtcaaagtttccggctacacaatccatgaatacacaggcaatctcggtggctgggttgacaggtgggttc  
agtacgacggatgggttaaaactgacggcacctctcacgatccagccaacggatattacggctactcagctctggagctacgcaggcgtcggt  
ga

SEQ ID NO: 112

Met Pro Ala Phe Lys Ser Lys Val Met His Met Lys Leu Lys Tyr Leu Ala Leu Val Leu Leu Ala Val  
Ala Ser Ile Gly Leu Leu Ser Thr Pro Val Gly Ala Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile  
Met Gln Ala Phe Tyr Trp Asp Val Pro Thr Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro  
Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser  
Met Gly Tyr Asp Pro Tyr Asp Phe Phe Asp Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr  
Arg Phe Gly Ser Lys Glu Glu Leu Val Asn Met Ile Asn Thr Ala His Ser Tyr Gly Ile Lys Val Ile  
Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp Leu Glu Trp Asn Pro Phe Val Asn Asn Tyr Thr  
Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu  
Val Lys Cys Cys Asp Glu Gly Thr Phe Gly Asp Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln  
Tyr Trp Leu Trp Ala Ser Asn Glu Ser Tyr Ala Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg  
Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val Asn Asp Trp Leu Ser Trp Trp Gly Gly Trp Ala  
Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Leu Leu Asn Trp Ala Tyr Asn Ser Gly Ala Lys Val  
Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala Phe Asp Asn Thr Asn Ile Pro Ala Leu Val Tyr  
Ala Leu Gln Asn Gly Gly Thr Val Val Ser Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His  
Asp Thr Asp Ile Ile Trp Asn Lys Tyr Pro Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile  
Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys Asp Lys Leu Asn Asn Leu Ile Trp Ile His Glu His  
Leu Ala Gly Gly Ser Thr Lys Ile Leu Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Met Arg Glu Gly Tyr  
Gly Ser Lys Pro Gly Leu Ile Thr Tyr Ile Asn Leu Gly Asn Asp Trp Ala Glu Arg Trp Val Asn Val  
Gly Ser Lys Phe Ala Gly Tyr Thr Ile His Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Arg Trp  
Val Gln Tyr Asp Gly Trp Val Lys Leu Thr Ala Pro Pro His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr  
Ser Val Trp Ser Tyr Ala Gly Val Gly

SEQ ID NO: 113

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ccgatcttaaaagcgagaaaacagtatgctacggagacagcatgattatttcgaccacatgacattgtcggctggacaagggaaggcgac  
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gacatggcatgacattaccggaaccgttcggagccggtgtcatcaattcggaggctggggagagtccacgtaaaccggcgggtcggttca  
atttatgttcaaagatag

Met Lys Gln Gln Lys Arg Leu Tyr Ala Arg Leu Leu Thr Leu Leu Phe Ala Leu Ile Phe Leu Leu Pro  
His Ser Ala Ala Ala Ala Ala Asn Leu Asn Gly Thr Leu Met Gln Tyr Phe Glu Trp Tyr Met Pro Asn  
Asp Gly Gln His Trp Lys Arg Leu Gln Asn Asp Ser Ala Tyr Leu Ala Glu His Gly Ile Thr Ala Val  
Trp Ile Pro Pro Ala Tyr Lys Gly Thr Ser Gln Ala Asp Val Gly Tyr Gly Ala Tyr Asp Leu Tyr Asp  
Leu Gly Glu Phe His Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr Lys Gly Glu Leu Gln Ser Ala  
Ile Lys Ser Leu His Ser Arg Asp Ile Asn Val Tyr Gly Asp Val Val Ile Asn His Lys Gly Gly Ala  
Asp Ala Thr Glu Asp Val Thr Ala Val Glu Val Asp Pro Ala Asp Arg Asn Arg Val Ile Ser Gly Glu  
His Arg Ile Lys Ala Trp Thr His Phe His Phe Pro Gly Arg Gly Ser Thr Tyr Ser Asp Phe Lys Trp  
His Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg Ile Tyr Lys Phe Gln  
Gly Lys Ala Trp Asp Trp Glu Val Ser Asn Glu Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Ile  
Asp Tyr Asp His Pro Asp Val Ala Ala Glu Ile Lys Arg Trp Gly Thr Trp Tyr Ala Asn Glu Leu Gln  
Leu Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Phe Ser Phe Leu Arg Asp Trp Val Asn His  
Val Arg Glu Lys Thr Gly Lys Glu Met Phe Thr Val Ala Glu Tyr Trp Gln Asn Asp Leu Gly Ala  
Leu Glu Asn Tyr Leu Asn Lys Thr Asn Phe Asn His Ser Val Phe Asp Val Pro Leu His Tyr Gln Phe  
His Ala Ala Ser Thr Gln Gly Gly Gly Tyr Asp Met Arg Lys Leu Leu Asn Gly Thr Val Val Ser Lys  
His Pro Leu Lys Ala Val Thr Phe Val Asp Asn His Asp Thr Gln Pro Gly Gln Ser Leu Glu Ser Thr  
Val Gln Thr Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Glu Ser Gly Tyr Pro Gln Val  
Phe Tyr Gly Asp Met Tyr Gly Thr Lys Gly Asp Ser Gln Arg Glu Ile Pro Ala Leu Lys His Lys Ile  
Glu Pro Ile Leu Lys Ala Arg Lys Gln Tyr Ala Tyr Gly Ala Gln His Asp Tyr Phe Asp His His Asp  
Ile Val Gly Trp Thr Arg Glu Gly Asp Ser Ser Val Ala Asn Ser Gly Leu Ala Ala Leu Ile Thr Asp  
Gly Pro Gly Gly Ala Lys Arg Met Tyr Val Gly Arg Gln Asn Ala Gly Glu Thr Trp His Asp Ile Thr  
Gly Asn Arg Ser Glu Pro Val Val Ile Asn Ser Glu Gly Trp Gly Glu Phe His Val Asn Gly Gly Ser  
Val Ser Ile Tyr Val Gln Arg

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ccggagtgggaagatctcatcaccatcaccatcactaa

Met Ala Lys Tyr Ser Glu Leu Glu Gln Gly Gly Val Ile Met Gln Ala Phe Tyr Trp Asp Val Pro Glu Gly Gly Ile Tyr Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile Ser Ala Ile Trp Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Ala Tyr Ser Met Gly Tyr Asp Pro Tyr Asp Tyr Phe Asp Leu Gly Glu Phe Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Glu Glu Leu Val Asn Met

Figure 16 (cont.)

Ile Ser Thr Ala His Gln Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg Ala Gly Gly Asp  
Leu Glu Trp Asn Pro Tyr Val Gly Asp Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala Ser Gly Lys Tyr  
Lys Ala His Tyr Met Asp Phe His Pro Asn Asn Tyr Ser Thr Ser Asp Glu Gly Thr Phe Gly Gly Phe  
Pro Asp Ile Asp His Leu Val Pro Phe Asn Gln Tyr Trp Leu Trp Ala Ser Asn Glu Ser Tyr Ala Ala  
Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly Ala Trp Val Val  
Lys Asp Trp Leu Ser Gln Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn Val Asp Ala Leu  
Leu Asn Trp Ala Tyr Ser Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys Met Asp Glu Ala  
Phe Asp Asn Lys Asn Ile Pro Ala Leu Val Tyr Ala Ile Gln Asn Gly Glu Thr Val Val Ser Arg Asp  
Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asn Ile Ile Trp Asn Lys Tyr Pro Ala Tyr  
Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu Trp Leu Asn Lys  
Asp Lys Leu Asn Asn Leu Ile Trp Ile His Glu His Leu Ala Gly Gly Ser Thr Lys Ile Leu Tyr Tyr  
Asp Asp Asp Glu Leu Ile Phe Met Arg Glu Gly Tyr Gly Asp Arg Pro Gly Leu Ile Thr Tyr Ile Asn  
Leu Gly Ser Asp Trp Ala Glu Arg Trp Val Asn Val Gly Ser Lys Phe Ala Gly Tyr Thr Ile His Glu  
Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Arg Tyr Val Gln Tyr Asp Gly Trp Val Lys Leu Thr Ala  
Pro Pro His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Ala Gly Val Gly Arg Ser  
His His His His His His

SEQ ID NO: 117

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ccaaggccgatcctcaaggtcgccggctacaagcaggtgctcctcaccgcctctgaagtcctcgggcaacgagtggtgggctcgttacc  
agccccaggtatcgccctggtcgacaccccccttggcaacaagcaggtatctggagcagctgatcgccgcgatgcagaccggggcattgc  
cgtctacgcggagctggtgctcaaccacatggccaacgaaagctggaagcgcagcgacctaactaccccggcagcgagctgctgcaaag  
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gtggctggcatcaacaagtgcgactacgagcaggagtactggctcgataccgccagattcgagatgaactggtatcgcaactaccgggatgtg  
ctcgaccagaatccggtggtcaactgcagagccagtgggtaaggctgaccatccggcccgccgcccaggaatgtgctgcaggagtga

SEQ ID NO: 118

Met Arg Val Phe Leu Val Val Pro Lys Leu Ser Arg Pro Phe Gln Ala Glu Ser Gln Gln Gln Asp Arg  
Asp Ile Thr Met Lys His Thr Ala Gly Met Leu Ala Ile Ala Gly Met Leu Ile Ala Pro Leu Ala His  
Ala Asp Val Ile Leu His Ala Phe Asn Trp Lys Tyr Ser Glu Val Thr Ala Lys Ala Asp Leu Ile Lys  
Ala Ala Gly Tyr Lys Gln Val Leu Ile Ser Pro Pro Leu Lys Ser Ser Gly Asn Glu Trp Trp Ala Arg  
Tyr Gln Pro Gln Asp Leu Arg Leu Val Asp Thr Pro Leu Gly Asn Lys Gln Asp Leu Glu Gln Leu Ile  
Ala Ala Met Gln Thr Arg Gly Ile Ala Val Tyr Ala Asp Val Val Leu Asn His Met Ala Asn Glu Ser  
Trp Lys Arg Ser Asp Leu Asn Tyr Pro Gly Ser Glu Leu Leu Gln Ser Tyr Ala Gly Asn Pro Ala Tyr  
Phe Glu Arg Gln Lys Leu Phe Gly Asp Leu Gly Gln Asn Phe Leu Ala Gly Gln Asp Phe His Pro  
Glu Gly Cys Ile Thr Asp Trp Asn Asn Pro Gly His Val Gln Tyr Trp Arg Leu Cys Gly Gly Ala Gly  
Asp Lys Gly Leu Pro Asp Leu Asp Pro Asn Asn Trp Val Val Asn Gln Gln Gln Ala Tyr Leu Gln  
Ala Leu Lys Gly Met Gly Ile Lys Gly Phe Arg Val Asp Ala Val Lys His Met Ser Asp Tyr Gln Ile  
Asn Ala Val Phe Thr Pro Glu Ile Lys Gln Gly Met His Val Phe Gly Glu Val Ile Thr Thr Gly Gly  
Ala Gly Asn Ser Asp Tyr Glu Asn Phe Leu Lys Pro Tyr Leu Asp Ser Ser Gly Gln Gly Ala Tyr Asp  
Phe Pro Leu Phe Ala Ser Leu Arg Gly Ala Leu Gly Tyr Gly Gly Ser Met Asn Leu Leu Ala Asp Pro

Figure 16 (cont.)

Gly Ala Tyr Gly Gln Ala Leu Pro Gly Ser Arg Ala Val Thr Phe Ala Ile Thr His Asp Ile Pro Thr  
Asn Asp Gly Phe Arg Tyr Gln Ile Leu Asn Gln Thr Asp Glu Arg Leu Ala Tyr Ala Tyr Leu Leu Gly  
Arg Asp Gly Gly Ser Pro Leu Val Tyr Ser Asp His Gly Glu Thr Arg Asp Lys Asp Gly Leu Arg Trp  
Gln Asp Tyr Tyr Leu Arg Thr Asp Leu Lys Gly Met Ile Arg Phe His Asn Thr Val Gln Gly Gln Pro  
Met Gln Leu Ile Gly Ser Asn Asp Cys Phe Val Leu Phe Lys Arg Gly Lys Gln Gly Val Val Gly Ile  
Asn Lys Cys Asp Tyr Glu Gln Glu Tyr Trp Leu Asp Thr Ala Arg Phe Glu Met Asn Trp Tyr Arg  
Asn Tyr Arg Asp Val Leu Asp Gln Asn Ala Val Val Asn Val Gln Ser Gln Trp Val Arg Leu Thr Ile  
Pro Ala Arg Gly Ala Arg Met Trp Leu Gln Glu

SEQ ID NO: 119

atgcaaacgtttgattcttattttactcaagaaaggatgggtgtgcatgaattatttgaaaaaagtggtgtattacgctatcgctaccttaa  
tcattctcttcttacaccttttcaacagcacaagctaatactgcacctgttaacggaacaatgatgcaataattcgaatgggacttacctaataatgatgg  
gacgctttggacgaaagtaaaaaatgaagctaccaatctttctcactaggtatcacagcactatggctccctccagcatataaaggaacgagcc  
aaagcgatgtcggatagcgtgtttacgatttatagacctggggaatttaataaaaaaggacgatccgaacgaaatacgaacaaaaacaca  
atatattcaagccattcaaaactgccccagccgagggatgcaagtatatgcggatgttgtatttaatacgaaggcaggggctgacagtacagaatt  
tgtcgatgcagttgaggttaaaccttctaatacgaaatcaagaacacatctggcacatatcaaatcaagcatggacaaaatttgattttctggtcgtg  
gaaacacatactccagcttcaaatggcgtgtgtaccattttgatgttacggattgggacgaaagtgtaaaattaaatcgtatttacaattccgcgg  
tacaggaaaagcgtgggactgggaagtcgatacagaaaacggaaactatgattatttaattgttcgctgatttagatatggatcacctgaggtgt  
gacagaattaaaaactggggaacgtgtgtacgtcaatactacaaatcatgatggattccgctagatgccgtaaaacataataacagcttttc  
cctgactggctaacatatgtacgtaatacaacaggaaaaaattatttgcctgtggggaattttggagctatgacgtcaataagctgcataattacat  
tacaaaaacaaatgggtcgatgtcattatttgatgcaccttgcataacaactttataccgcttccaaatcgagtggatatttgacatgcgttattat  
tgaataatacattaatgaaagatcaaccttcactcgtgtaacacttgcgataaccagacacgcaaccagggcaatctttacagtcaggtgcg  
aaccttggtttaaacagcttgccttacgctttattttaacaagacaagaagggtatccttgcgtattttacggtgattattatggaatccctaaatacaat  
atccccggggttaaaaagtaaaatcgaccgcttttaattgtcgtcgtgattacgcttatggaacacaacgtgattacattgatcatcaagacattat  
cggatggacacgagaaggcattgatgcaaaaccgaactctggactggcggctttaattaccgacggctcgtggaagtaaatggatgtatgtc  
ggtaaaaagcatgccgggaaagtattttatgatttaactggaatcgaagtacacagtaacgattaatgcggatggttggggagaatttaagta  
aacggaggatccgtctcaatttgggtggctaaacgtcaaacgtcacatttacagtaataacgccacaacaacagcggacaaaacgtatatg  
ttgcggcaacattccagagctaggcaattgtcgcacgggttaa

SEQ ID NO: 120

Met Gln Thr Phe Ala Phe Leu Phe Tyr Ser Lys Lys Gly Trp Val Cys Met Asn Tyr Leu Lys Lys Val  
Trp Leu Tyr Tyr Ala Ile Val Ala Thr Leu Ile Ile Ser Phe Leu Thr Pro Phe Ser Thr Ala Gln Ala Asn  
Thr Ala Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu Trp Asp Leu Pro Asn Asp Gly Thr Leu  
Trp Thr Lys Val Lys Asn Glu Ala Thr Asn Leu Ser Ser Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro  
Ala Tyr Lys Gly Thr Ser Gln Ser Asp Val Gly Tyr Gly Val Tyr Asp Leu Tyr Asp Leu Gly Glu Phe  
Asn Gln Lys Gly Thr Ile Arg Thr Lys Tyr Gly Thr Lys Thr Gln Tyr Ile Gln Ala Ile Gln Thr Ala  
Gln Ala Ala Gly Met Gln Val Tyr Ala Asp Val Val Phe Asn His Lys Ala Gly Ala Asp Ser Thr Glu  
Phe Val Asp Ala Val Glu Val Asn Pro Ser Asn Arg Asn Gln Glu Thr Ser Gly Thr Tyr Gln Ile Gln  
Ala Trp Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr His  
Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg Ile Tyr Lys Phe Arg Gly Thr Gly Lys  
Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn Tyr Asp Tyr Leu Met Phe Ala Asp Leu Asp  
Met Asp His Pro Glu Val Val Thr Glu Leu Lys Asn Trp Gly Thr Trp Tyr Val Asn Thr Thr Asn Ile  
Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Tyr Ser Phe Phe Pro Asp Trp Leu Thr Tyr Val  
Arg Asn Gln Thr Gly Lys Asn Leu Phe Ala Val Gly Glu Phe Trp Ser Tyr Asp Val Asn Lys Leu His  
Asn Tyr Ile Thr Lys Thr Asn Gly Ser Met Ser Leu Phe Asp Ala Pro Leu His Asn Asn Phe Tyr Thr  
Ala Ser Lys Ser Ser Gly Tyr Phe Asp Met Arg Tyr Leu Leu Asn Asn Thr Leu Met Lys Asp Gln  
Pro Ser Leu Ala Val Thr Leu Val Asp Asn His Asp Thr Gln Pro Gly Gln Ser Leu Gln Ser Trp Val  
Glu Pro Trp Phe Lys Gln Leu Ala Tyr Ala Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe  
Tyr Gly Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro Gly Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile  
Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln Arg Asp Tyr Ile Asp His Gln Asp Ile Ile Gly Trp Thr

Figure 16 (cont.)

Arg Glu Gly Ile Asp Ala Lys Pro Asn Ser Gly Leu Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser  
Lys Trp Met Tyr Val Gly Lys Lys His Ala Gly Lys Val Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp  
Thr Val Thr Ile Asn Ala Asp Gly Trp Gly Glu Phe Lys Val Asn Gly Gly Ser Val Ser Ile Trp Val  
Ala Lys Thr Ser Asn Val Thr Phe Thr Val Asn Asn Ala Thr Thr Thr Ser Gly Gln Asn Val Tyr Val  
Val Gly Asn Ile Pro Glu Leu Gly Asn Cys Arg Thr Gly

SEQ ID NO: 121

atgctcgcctgtcgtcggcggtgcggcatcgacgcggggcccgacaggccctcgcgtcgtggagccgctgccgcagcgcggccacgcttc  
cgcaggagtagccgcgccagcggccacgcggccgcccggcgacgtgttcgtgcacctgttcgagtggaagtggccggacatcgccggaggaa  
gcgagaacgtgctggggccggcggtgtacgaggcggtgcaggtgtcggccgcaggagcacctggtgcagcagggggcgccgtggtg  
gcagcggtagccagccggtgagctactcgggtggcgctgagccgcagcggcgacggcggtggagttcagcaacatgatcagccggtgcaaggc  
cgccggcggtggacatctacgtggacgccgtatcaaccacatgacggccggtgcggggacggggagcaacggcaccgcctacaccaagta  
caactacccggcctgtacgcgcagcgggactttcacccgcagtgcgcgggtggcgactacaccagcggcccaacgtgcaggactgcga  
actgctggggctggctgacctgaacaccggcgcgccggcggtgcagcagaagatcgccggactacctggtcgcgtggcgcggtggcggt  
ggcggggttttcgacgcagccgccaagcacatccagccggtggaactggacgccatcgtggaccgcgtgaaccagacgctggcgccgga  
ggggcgcccgcttcctactggttcgcccaggtgatcgacaacggcgcgagggggtgcggcgcgagcactactacggcctgggatacgg  
caccggcgccgcccggacatcacggagttccgctacaagggcggtggcgacaaagtctcgggcagcggcgccagcggctggtggacc  
tgaagaactctcggcggtgacgtggaacctgatccgctcggaacggcgctcgtctttcggagaaccacgatacgcagcgcggcgccggc  
atcggtaccgcgatggcacggcggttcggctggccaacgtgtggatgctggcgagccgtacggctatccgtcggtgatgccagctacgc  
ctttgaccgcacctccccctttggccgcgacgccggcccgccctccgaggacggcgacgaaggacgtgacgtgcgcccacgctgga  
gacggcggtgctgggacacctgggtgtgcgagcaccgcgaccccgctcattcagcggatggtgggctttcgccgcgcgatggcgggcacgga  
cctgaaccgctggtgggacaacggcggaacgccattgccttttcgcgcggggaccggggttcgctgccatcagccgcgagccgaagggtg  
accatggcgccgctgccagcggactgtccccggcacctactgcgacgtgctgaccggcggaaggtgggcaacgcctgcgcgggaac  
cagcgtgacggctgactctcaggcggtggtgcagctgagcatcgtcgagaactcggctctggtgatccacctcggggccaagctgtaacggc  
gcgctggcggtggtgcggagggg

SEQ ID NO: 122

Met Leu Ala Leu Ser Leu Gly Gly Cys Gly Ile Asp Ala Gly Pro Thr Gly Pro Arg Val Val Glu Pro  
Leu Pro Gln Arg Pro Thr Leu Pro Gln Glu Tyr Arg Ala Ser Gly His Ala Ala Ala Gly Asp Val Phe  
Val His Leu Phe Glu Trp Lys Trp Pro Asp Ile Ala Glu Glu Cys Glu Asn Val Leu Gly Pro Ala Gly  
Tyr Glu Ala Val Gln Val Ser Pro Pro Gln Glu His Leu Val Gln Gln Gly Ala Pro Trp Trp Gln Arg  
Tyr Gln Pro Val Ser Tyr Ser Val Ala Leu Ser Arg Ser Gly Thr Gly Val Glu Phe Ser Asn Met Ile  
Ser Arg Cys Lys Ala Ala Gly Val Asp Ile Tyr Val Asp Ala Val Ile Asn His Met Thr Ala Gly Ala  
Gly Thr Gly Ser Asn Gly Thr Ala Tyr Thr Lys Tyr Asn Tyr Pro Gly Leu Tyr Ala Gln Ala Asp Phe  
His Pro Gln Cys Ala Val Gly Asp Tyr Thr Ser Ala Ala Asn Val Gln Asp Cys Glu Leu Leu Gly Leu  
Ala Asp Leu Asn Thr Gly Ala Ala Gly Val Gln Gln Lys Ile Ala Asp Tyr Leu Val Ser Leu Ala Arg  
Leu Gly Val Ala Gly Phe Arg Ile Asp Ala Ala Lys His Ile Gln Pro Val Glu Leu Asp Ala Ile Val  
Asp Arg Val Asn Gln Thr Leu Ala Ala Glu Gly Arg Pro Leu Pro Tyr Trp Phe Ala Glu Val Ile Asp  
Asn Gly Gly Glu Gly Val Arg Arg Glu His Tyr Tyr Gly Leu Gly Tyr Gly Thr Gly Gly Ala Ala Asp  
Ile Thr Glu Phe Arg Tyr Lys Gly Val Gly Asp Lys Phe Leu Gly Ser Gly Gly Gln Arg Leu Val Asp  
Leu Lys Asn Phe Ser Ala Val Thr Trp Asn Leu Met Pro Ser Asp Lys Ala Val Val Phe Leu Glu Asn  
His Asp Thr Gln Arg Gly Gly Gly Ile Gly Tyr Arg Asp Gly Thr Ala Phe Arg Leu Ala Asn Val Trp  
Met Leu Ala Gln Pro Tyr Gly Tyr Pro Ser Val Met Ser Ser Tyr Ala Phe Asp Arg Thr Ser Pro Phe  
Gly Arg Asp Ala Gly Pro Pro Ser Glu Asp Gly Ala Thr Lys Asp Val Thr Cys Ala Pro Thr Leu Glu  
Thr Ala Val Leu Gly Thr Trp Val Cys Glu His Arg Asp Pro Val Ile Gln Arg Met Val Gly Phe Arg  
Arg Ala Met Ala Gly Thr Asp Leu Asn Arg Trp Trp Asp Asn Gly Gly Asn Ala Ile Ala Phe Ser Arg  
Gly Asp Arg Gly Phe Val Ala Ile Ser Arg Glu Pro Lys Val Thr Met Ala Ala Val Pro Ser Gly Leu  
Ser Pro Gly Thr Tyr Cys Asp Val Leu Thr Gly Gly Lys Val Gly Asn Ala Cys Ala Gly Thr Ser Val  
Thr Val Asp Ser Gln Gly Val Val Gln Leu Ser Ile Val Glu Asn Ser Ala Leu Val Ile His Leu Gly  
Ala Lys Leu Arg Arg Ala Gly Gly Cys Ala Glu

Figure 16 (cont.)

SEQ ID NO: 123

atgccccaggccattgcactttttcacgttggacgtgttcggcttaatcggcgttttctgcttggctctgcttttctgtcccacccgggcaatcc  
aggccagacaaccccgccccgtaccgttatggttcacctcttcgagtggaaatggaccgacatcgctaaagaatgcgagaatttctcggac  
cgaaaggctttgcccgaatccaggtatgccgccccaggagcatgtccaggggtcgcaatggtggaccgcgtatcagccggtcagctacaag  
atcgagagccgctccggcaccgggcccaggttcgccaatatggtctcgcgtgcaaagccgtcgggggtgatatctatgtcgtatgccgtgatc  
aaccatatgacgactgtcggctccggcactgggtatggctggatcgacctacaccagctacacctatccggggctgtatcagacccaggacttcc  
accactgcggggcgcaatggcaacgatgatatcagcagctacggcgatcgtcgggaagtacaaaactcggaactgctaacctagccgacctc  
aacaccggcgctgagtatgtccgggtaaacctcggcgctatatgaacgatctcgcggcctggggcgtcggcgatttcggatcgatccggcc  
aagccatggataccaacgacatcaacaatctcgttggccgcctgccccaacgcgcctacatctaccaggaagtgtatcgaccagggcgccga  
ggcaattaccggcggaatacttcagaatggcgatgtgaccgaggtcaagtacagccgcgagatctcgcgcagtgttcaaaaccggccagct  
gacctatagccagttcggcactgcctgggggttcatgtccagcgacctggcagtagttttaccgataaccacgacaaccagcgccgtca  
cggcgggcgccggcgatgtcttgacctacaagatggccagctgtacacctgggcaatatcttcgagctagcctggccgtatggctaccaca  
ggatcatgtcgagctacacgttcagcaacggcgaccagggggcgccatcgaccaatgtgtacgcaaccacaacgctgattgtggaacggcc  
gctgggtctgtgagcaccgctggcgaggaatcgccaacatggtcgcgttcgcaactacaccgccccgaccttcagcaccagcaactggtgg  
agcaacggcaacaaccagatcgctttcagccgcccggaccctgggctttgtggcgatcaatcgggaaggtggcagcctgaaccgaccttcca  
aaccggcctgcccgtcggcacctactgcgatgtcattcacggcgatttcaatgccagcgccggcacctgttccggccaactatcgctgtcaac  
ggctccggacaggcaaccatcacggtaacgcgatggacgcgggtggcgatctacggcgagccaggtcgcgacctccggccagtgtcaac  
gtgacattcaacgaaaacgccacgaccacctgggggcagaatgtgtatatcgtcggcaacgtcggcgccctgggcagctggaacgcaggca  
gcgcggttactctctccgtaactacccaatctggagcaagaccatcgccctgccagccaacaccgcattgagtacaagtacatcaaaaa  
ggatggcgccgggcaatgtggtgtgggaaagcggcgccaaccgcgtctttaccacccccggcagcggcagtgccacgcgcaacgatacctg  
gaaatag

SEQ ID NO: 124

Met Pro Gln Ala Ile Arg Thr Phe Ser Arg Trp Thr Leu Phe Gly Leu Ile Gly Val Phe Leu Leu Gly  
Leu Val Phe Ser Val Pro Pro Arg Ala Ile Gln Ala Gln Thr Thr Pro Ala Arg Thr Val Met Val His  
Leu Phe Glu Trp Lys Trp Thr Asp Ile Ala Lys Glu Cys Glu Asn Phe Leu Gly Pro Lys Gly Phe Ala  
Ala Ile Gln Val Ser Pro Pro Gln Glu His Val Gln Gly Ser Gln Trp Trp Thr Arg Tyr Gln Pro Val Ser  
Tyr Lys Ile Glu Ser Arg Ser Gly Thr Arg Ala Glu Phe Ala Asn Met Val Ser Arg Cys Lys Ala Val  
Gly Val Asp Ile Tyr Val Asp Ala Val Ile Asn His Met Thr Thr Val Gly Ser Gly Thr Gly Met Ala  
Gly Ser Thr Tyr Thr Ser Tyr Thr Tyr Pro Gly Leu Tyr Gln Thr Gln Asp Phe His His Cys Gly Arg  
Asn Gly Asn Asp Asp Ile Ser Ser Tyr Gly Asp Arg Trp Glu Val Gln Asn Cys Glu Leu Leu Asn  
Leu Ala Asp Leu Asn Thr Gly Ala Glu Tyr Val Arg Gly Lys Leu Ala Ala Tyr Met Asn Asp Leu  
Arg Gly Leu Gly Val Ala Gly Phe Arg Ile Asp Ala Ala Lys His Met Asp Thr Asn Asp Ile Asn Asn  
Ile Val Gly Arg Leu Pro Asn Ala Pro Tyr Ile Tyr Gln Glu Val Ile Asp Gln Gly Gly Glu Pro Ile Thr  
Ala Gly Glu Tyr Phe Gln Asn Gly Asp Val Thr Glu Phe Lys Tyr Ser Arg Glu Ile Ser Arg Met Phe  
Lys Thr Gly Gln Leu Thr His Met Ser Gln Phe Gly Thr Ala Trp Gly Phe Met Ser Ser Asp Leu Ala  
Val Val Phe Thr Asp Asn His Asp Asn Gln Arg Gly His Gly Gly Ala Gly Asp Val Leu Thr Tyr  
Lys Asp Gly Gln Leu Tyr Thr Leu Gly Asn Ile Phe Glu Leu Ala Trp Pro Tyr Gly Tyr Pro Gln Val  
Met Ser Ser Tyr Thr Phe Ser Asn Gly Asp Gln Gly Pro Pro Ser Thr Asn Val Tyr Ala Thr Thr Thr  
Pro Asp Cys Gly Asn Gly Arg Trp Val Cys Glu His Arg Trp Arg Gly Ile Ala Asn Met Val Ala Phe  
Arg Asn Tyr Thr Ala Pro Thr Phe Ser Thr Ser Asn Trp Trp Ser Asn Gly Asn Asn Gln Ile Ala Phe  
Ser Arg Gly Thr Leu Gly Phe Val Ala Ile Asn Arg Glu Gly Gly Ser Leu Asn Arg Thr Phe Gln Thr  
Gly Leu Pro Val Gly Thr Tyr Cys Asp Val Ile His Gly Asp Phe Asn Ala Ser Ala Gly Thr Cys Ser  
Gly Pro Thr Ile Ala Val Asn Gly Ser Gly Gln Ala Thr Ile Thr Val Asn Ala Met Asp Ala Val Ala Ile  
Tyr Gly Gly Ala Arg Leu Ala Thr Pro Ala Ser Val Asn Val Thr Phe Asn Glu Asn Ala Thr Thr Thr  
Trp Gly Gln Asn Val Tyr Ile Val Gly Asn Val Ala Ala Leu Gly Ser Trp Asn Ala Gly Ser Ala Val  
Leu Leu Ser Ser Ala Asn Tyr Pro Ile Trp Ser Lys Thr Ile Ala Leu Pro Ala Asn Thr Ala Ile Glu Tyr  
Lys Tyr Ile Lys Lys Asp Gly Ala Gly Asn Val Val Trp Glu Ser Gly Ala Asn Arg Val Phe Thr Thr  
Pro Gly Ser Gly Ser Ala Thr Arg Asn Asp Thr Trp Lys

Figure 16 (cont.)

SEQ ID NO: 125

gtggtgcacatgaagtgaagtaccttgccttagtttgggtgtggttcgataggcctactctgactccagtgggtgctgccaagtactccg  
aactcgaagagggcggtgttataatgcaggccttctactgggatgttcccgagggggaatctggtgggacaccataagacagaaaatccc  
gagtggtacgacgctggaatctcggcgatatggattcctccagctagcaaagggtggcggtgttattccatgggctacgatccctacgattt  
ctttgacctcggcgagtactatcagaagggaacagttgagacgcgcttcggctcaaaggaggaactggtgaacatgataaacaccgcacactc  
ctatggcataaagggtgatacgggacatagtcataaaccaccgcgcgggtggagaccttgagtgaacccctttgtaacaactatacttggaca  
gacttctccaaggctgcctccggtaatacacggccaactaccttgacttccacccaaacgaggtcaagtctgcgatgagggtacatttggta  
ctttccggacatcggccacgagaagagctgggatcagctactggctctgggcaagcaatgagagctacgcccgcatactccggagcataggga  
tcgatgcatggcgcttgcactacgtaaaaggttacggagcgtgggtgttaatgactggctcagctggtgggagggtggccggtggagagta  
ctgggacacgaacgttgatgcactcctaactgggcatacgcagcggtgccaaggctttgacttcccgcctactacaagatggacgaagcc  
tttgacaacaccaacatccccgcttggtttgcctccagaacggaggaacagctgttcccgcgatccctcaaggcagtaacttctgttgc  
aaccacgatacagataatctggaacaagtatccggcttatgcgttcacttacctatgagggacagcctgttatatttaccgcgactacgagg  
agtggctcaacaaggataagcttaacaaccttatctggatacagcagccttgcggagggaagtaaccaagatcctctactacgataacgatga  
gctaattatcatgagggagggtacgggagcaagccgggctcataacctacataaacctcggaaacgactgggcccagcgtgggtgaac  
gtcggctcaaaagtgttccggctacacaatccatgaatacacaggcaatctcgggtgggtgacaggtgggttcagtagtgatgggtta  
aactgacggcaccctctcatgatccagccaacggatattacggctactcagctggagctacgcaggcgtcggatga

SEQ ID NO: 126

Val Val His Met Lys Leu Lys Tyr Leu Ala Leu Val Leu Ala Val Ala Ser Ile Gly Leu Leu Ser  
Thr Pro Val Gly Ala Ala Lys Tyr Ser Glu Leu Glu Glu Gly Gly Val Ile Met Gln Ala Phe Tyr Trp  
Asp Val Pro Gly Gly Gly Ile Trp Trp Asp Thr Ile Arg Gln Lys Ile Pro Glu Trp Tyr Asp Ala Gly Ile  
Ser Ala Ile Trp Ile Pro Pro Ala Ser Lys Gly Met Gly Gly Gly Tyr Ser Met Gly Tyr Asp Pro Tyr  
Asp Phe Phe Asp Leu Gly Glu Tyr Tyr Gln Lys Gly Thr Val Glu Thr Arg Phe Gly Ser Lys Glu Glu  
Leu Val Asn Met Ile Asn Thr Ala His Ser Tyr Gly Ile Lys Val Ile Ala Asp Ile Val Ile Asn His Arg  
Ala Gly Gly Asp Leu Glu Trp Asn Pro Phe Val Asn Asn Tyr Thr Trp Thr Asp Phe Ser Lys Val Ala  
Ser Gly Lys Tyr Thr Ala Asn Tyr Leu Asp Phe His Pro Asn Glu Val Lys Cys Cys Asp Glu Gly Thr  
Phe Gly Asp Phe Pro Asp Ile Ala His Glu Lys Ser Trp Asp Gln Tyr Trp Leu Trp Ala Ser Asn Glu  
Ser Tyr Ala Ala Tyr Leu Arg Ser Ile Gly Ile Asp Ala Trp Arg Phe Asp Tyr Val Lys Gly Tyr Gly  
Ala Trp Val Val Asn Asp Trp Leu Ser Trp Trp Gly Gly Trp Ala Val Gly Glu Tyr Trp Asp Thr Asn  
Val Asp Ala Leu Leu Asn Trp Ala Tyr Asp Ser Gly Ala Lys Val Phe Asp Phe Pro Leu Tyr Tyr Lys  
Met Asp Glu Ala Phe Asp Asn Thr Asn Ile Pro Ala Leu Val Tyr Ala Leu Gln Asn Gly Gly Thr Val  
Val Ser Arg Asp Pro Phe Lys Ala Val Thr Phe Val Ala Asn His Asp Thr Asp Ile Ile Trp Asn Lys  
Tyr Pro Ala Tyr Ala Phe Ile Leu Thr Tyr Glu Gly Gln Pro Val Ile Phe Tyr Arg Asp Tyr Glu Glu  
Trp Leu Asn Lys Asp Lys Leu Asn Asn Leu Ile Trp Ile His Glu His Leu Ala Gly Gly Ser Thr Lys  
Ile Leu Tyr Tyr Asp Asn Asp Glu Leu Ile Phe Met Arg Glu Gly Tyr Gly Ser Lys Pro Gly Leu Ile  
Thr Tyr Ile Asn Leu Gly Asn Asp Trp Ala Glu Arg Trp Val Asn Val Gly Ser Lys Phe Ala Gly Tyr  
Thr Ile His Glu Tyr Thr Gly Asn Leu Gly Gly Trp Val Asp Arg Trp Val Gln Tyr Asp Gly Trp Val  
Lys Leu Thr Ala Pro Pro His Asp Pro Ala Asn Gly Tyr Tyr Gly Tyr Ser Val Trp Ser Tyr Ala Gly  
Val Gly

SEQ ID NO: 127

gtgtgcatgaattatttgaaaaaagtgtggtgtattacgctatcgtcgtacttcaatcatttcttctacgccctttcaactgcacaagccaacac  
tgcaccagtcaacggaacgatgatgcaatatttgaatgggattaccgaatgatggcacacttggacgaaagtaaaaaacgaagcaagcagt  
ctttcttcttaggtattactgcgttatggttaccacctgcatacaaaaggaacgagccaaggggatgctgggtatggcgtgtacgatttgtatgactt  
aggagaatttaatacaaaaagggacgattcgaacgaatacggaaacaaaacgcaatattacaagccattcaagcggcaaaaagcgctggcat  
gcaagtatacgtgatgtcgtatttaacacaaggcgggggcagatagtagaagaatgggttgacgcagtcgaagtgaatccttcaatcgaacc  
aagaaacatctggcacatatcaaatcaagcatggacaaaattgtattccctggccgtgggaacacatactcaagctttaaattggcgatggatc  
attttgacggtaggattgggatgaaagccgaaaactaaatcgtattacaatttcgtggcacaggaaaagcatgggattgggaagtagacaca  
gagaacggaaactatgactactaatgttctgatttagatatggatcacctgaaagtcgtgacagagctaaaaaactggggaacatggtacgtc

Figure 16 (cont.)

aatacagacaaatgctgatgggttcgcttagatgcagtaaagcatattaaatatagcttctccagattgggtaacacatgtgcgttcacaaacacg  
aaaaaatcttttgcagtaggagaattttggagctacgatgtcaataaactgcataactacattacaaaaacaagtgaaccatgtcgttatttgatg  
cgccacttcataacaacttttactgcttcaaaatctagcgggtattttgacatgcgctattttgtaataatacgttgatgaaagaccagccttctt  
tgcgggtcacactcgttgataatcatgacacgcaaccgggacaatctttacaatcatgggtagagccttggttaagccgcttgcttatgcctttattt  
gacaagacaagaaggataccttgctgtattttacggcgactattacggcatccctaaatacaacattccgggattgaaaagtaaaatcgatccgct  
tctcattgcccgtagagactacgcatacggaacacaacgtgattatattgaccatcaagacattattggatggacacgggaaggaattgactcaa  
aaccgaactctggacttgcggcttaattactgacggccctgggtggaagtaaatggatgtatgtaggtaaaaagcatgctggaaaagtgttttacg  
atctcactggaaatcgaaagcagacggtaacgattaatgcagacggctggggagagttaaagtaaacgggtggctccgtttccatttgggttgc  
aaaacatcacaaagtcacgtttaccgtcaacaatgcgacaacgataagcggacaaaatgtgtatgtcgttgtaacattccagagctcggaattg  
gaacacagcaaacgcaatcaaaatgaccccatcttctatccaacgttgaaagcaaccattgcttccacaaggaaaagccattgaatttaaatt  
tattaaaaagaccaatcgggaaatgtgttgggaaagcattccaaccgaacataccggtccattttatcaacaggctcatatacagctagt  
ggaatgtacctta

SEQ ID NO: 128

Val Cys Met Asn Tyr Leu Lys Lys Val Trp Leu Tyr Tyr Ala Ile Val Ala Thr Leu Ile Ile Tyr Phe  
Leu Thr Pro Phe Ser Thr Ala Gln Ala Asn Thr Ala Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu  
Trp Asp Leu Pro Asn Asp Gly Thr Leu Trp Thr Lys Val Lys Asn Glu Ala Ser Ser Leu Ser Ser Leu  
Gly Ile Thr Ala Leu Trp Leu Pro Ala Tyr Lys Gly Thr Ser Gln Gly Asp Val Gly Tyr Gly Val  
Tyr Asp Leu Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Ile Arg Thr Lys Tyr Gly Thr Lys Thr  
Gln Tyr Leu Gln Ala Ile Gln Ala Ala Lys Ser Ala Gly Met Gln Val Tyr Ala Asp Val Val Phe Asn  
His Lys Ala Gly Ala Asp Ser Thr Glu Trp Val Asp Ala Val Glu Val Asn Pro Ser Asn Arg Asn Gln  
Glu Thr Ser Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn Thr Tyr  
Ser Ser Phe Lys Trp Arg Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg  
Ile Tyr Lys Phe Arg Gly Thr Gly Lys Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn Tyr Asp  
Tyr Leu Met Phe Ala Asp Leu Asp Met Asp His Pro Glu Val Val Thr Glu Leu Lys Asn Trp Gly  
Thr Trp Tyr Val Asn Thr Thr Asn Val Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Tyr Ser  
Phe Phe Pro Asp Trp Leu Thr His Val Arg Ser Gln Thr Arg Lys Asn Leu Phe Ala Val Gly Glu Phe  
Trp Ser Tyr Asp Val Asn Lys Leu His Asn Tyr Ile Thr Lys Thr Ser Gly Thr Met Ser Leu Phe Asp  
Ala Pro Leu His Asn Asn Phe Tyr Thr Ala Ser Lys Ser Ser Gly Tyr Phe Asp Met Arg Tyr Leu Leu  
Asn Asn Thr Leu Met Lys Asp Gln Pro Ser Leu Ala Val Thr Leu Val Asp Asn His Asp Thr Gln  
Pro Gly Gln Ser Leu Gln Ser Trp Val Glu Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr  
Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro Gly  
Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln Arg Asp Tyr  
Ile Asp His Gln Asp Ile Ile Gly Trp Thr Arg Glu Gly Ile Asp Ser Lys Pro Asn Ser Gly Leu Ala Ala  
Leu Ile Thr Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Lys His Ala Gly Lys Val Phe  
Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn Ala Asp Gly Trp Gly Glu Phe Lys Val  
Asn Gly Gly Ser Val Ser Ile Trp Val Ala Lys Thr Ser Gln Val Thr Phe Thr Val Asn Asn Ala Thr  
Thr Ile Ser Gly Gln Asn Val Tyr Val Val Gly Asn Ile Pro Glu Leu Gly Asn Trp Asn Thr Ala Asn  
Ala Ile Lys Met Thr Pro Ser Ser Tyr Pro Thr Trp Lys Ala Thr Ile Ala Leu Pro Gln Gly Lys Ala Ile  
Glu Phe Lys Phe Ile Lys Lys Asp Gln Ser Gly Asn Val Val Trp Glu Ser Ile Pro Asn Arg Thr Tyr  
Thr Val Pro Phe Leu Ser Thr Gly Ser Tyr Thr Ala Ser Trp Asn Val Pro

SEQ ID NO: 129

ttgcgttgcgcgctggcagggacgggtgttgggtgcggcgcgtaatgcgctgccgcgacacccgcgtgaacaaaataatgaattattg  
aataggatgggggtgtcaagaatgacaaaatctcagagttgcggtgtcatggaagattttgttgggtgctgttggatggcttgggga  
tcttcgcgctcgcggcgctattgatgcaaggcttactgggacgccagtagccgggaccagtgattcgttggtggacgattggccaagcaag  
ccaacggtctaaaacgggcggggttcaccgccgtatggattcctccggtgtctaaaggggcttcagggggctattccaacgggtacgatccctt  
tgacgactatgatacgaagcaaggaccagaaaaggtaccgtggcgacgcgatgggggacgcgagaagaactgcaacgtgccgtggccgt  
gatgcgcgcgaacgggtcgtgatgtatgtggatctggtgctgaaccaccgcaacggggacgacgggaattggaatttcattacaagatgc  
gtacggcaaaagtggttacggcggttcaaaaggggtttacgattttaccccaactacaacattcaggatgccaatgttcccaacgaggattc

Figure 16 (cont.)

cagcttcgggcgcgatttagcccatgacaatccgtatgtggccgatggactgaaggctgcaggcgattggctgaccaaagccctcgatgttca  
gggatacgtctggattacgtgaaggcatcagctacaccttctgaaaagtattctgtcctatggggccatgaacggaaaatttgcgctcggtga  
gtactgggatgccaacgggatacgttgaactgggtggcgaacacggcgatggaaggcgggcccatgtgttgatttgcgttcgcgagg  
agctgaaaaacatgtgcaatcgggacgggtactacgacatgcgtcgattggaccacggggtctggtcggaatcaccctggaaggcggt  
gacgttttcgaaaacatgatacggatcggcacgaccccatctacaataacaagcatttggcgtatgcctacatcttgacgtcgaagggtatc  
cgacgggtgttctggaaggattactaccaatacggaatgaagccgatcatcgacaacctcatttggatccacgaacacattgcgtacggaacgac  
ccaagagcgttggaaagacgaagatgtctttgtgtatgagcggaccggaggcaagcggctattgggtggggttaacgacaatcgcgccacca  
gcaaacgggtcaccgtacagaccggttgggtgccaacgtggccttgacgactacaccggcaacggccccgatctccgtaccgacgcctac  
ggtcgggtaaccttgaccattcctgcaaacgggtacgtggcctattcgttccgggcatctccggatccttggcgggtcgagaaaaccgtgac  
gcaggagtttgcggggcgctccgacttgatattcgtccggcgataacacgcaatttgcaggctcggcggtatatacgcaaggcaaacaa  
gccggttacagcggaattgtattgggatgccaagactggacgacctccacgtcgattctcctagaagtgcgttcggcttcgggaacgctcatc  
acgacaaagaccgtgacccaattgtcgtccagggtaccgcggttcttcacgccttcggctaccggatggtacgtcttttccattcgaagctat  
aacacgccttcgacgaacccaagccggcctactggttaaaggtaacgtatacggcgccgaattgcttcagtaa

SEQ ID NO: 130

Met Arg Cys Arg Arg Gly Arg Asp Gly Cys Trp Cys Gly Arg Arg Asn Ala Leu Pro Arg His Pro  
Arg Glu Gln Asn Asn Met Asn Tyr Leu Asn Arg Met Gly Val Ser Arg Met Thr Lys Ser Arg Glu  
Leu Arg Cys Ser Trp Lys Val Phe Val Val Gly Cys Leu Leu Trp Met Ala Trp Gly Ser Ser Ala Ser  
Ala Gly Val Leu Met Gln Gly Phe Tyr Trp Asp Ala Ser Thr Gly Thr Ser Asp Ser Trp Trp Thr His  
Leu Ala Lys Gln Ala Asn Gly Leu Lys Arg Ala Gly Phe Thr Ala Val Trp Ile Pro Pro Val Leu Lys  
Gly Ala Ser Gly Gly Tyr Ser Asn Gly Tyr Asp Pro Phe Asp Asp Tyr Asp Ile Gly Ser Lys Asp Gln  
Lys Gly Thr Val Ala Thr Arg Trp Gly Thr Arg Glu Glu Leu Gln Arg Ala Val Ala Val Met Arg Ala  
Asn Gly Leu Asp Val Tyr Val Asp Leu Val Leu Asn His Arg Asn Gly Asp Asp Gly Asn Trp Asn  
Phe His Tyr Lys Asp Ala Tyr Gly Lys Val Gly Tyr Gly Arg Phe Gln Lys Gly Phe Tyr Asp Phe His  
Pro Asn Tyr Asn Ile Gln Asp Ala Asn Val Pro Asn Glu Asp Ser Ser Phe Gly Arg Asp Leu Ala His  
Asp Asn Pro Tyr Val Ala Asp Gly Leu Lys Ala Ala Gly Asp Trp Leu Thr Lys Ala Leu Asp Val  
Gln Gly Tyr Arg Leu Asp Tyr Val Lys Gly Ile Ser Tyr Thr Phe Leu Lys Ser Tyr Leu Ser Tyr Gly  
Ala Met Asn Gly Lys Phe Ala Val Gly Glu Tyr Trp Asp Ala Asn Arg Asp Thr Leu Asn Trp Trp  
Ala Asn Thr Ala Met Glu Gly Arg Ala His Val Phe Asp Phe Ala Leu Arg Glu Glu Leu Lys Asn  
Met Cys Asn Ala Asp Gly Tyr Tyr Asp Met Arg Arg Leu Asp His Ala Gly Leu Val Gly Ile Asp  
Pro Trp Lys Ala Val Thr Phe Val Glu Asn His Asp Thr Asp Arg His Asp Pro Ile Tyr Asn Asn Lys  
His Leu Ala Tyr Ala Tyr Ile Leu Thr Ser Glu Gly Tyr Pro Thr Val Phe Trp Lys Asp Tyr Tyr Gln  
Tyr Gly Met Lys Pro Ile Ile Asp Asn Leu Ile Trp Ile His Glu His Ile Ala Tyr Gly Thr Thr Gln Glu  
Arg Trp Lys Asp Glu Asp Val Phe Val Tyr Glu Arg Thr Gly Gly Lys Arg Leu Leu Val Gly Leu  
Asn Asp Asn Arg Ala Thr Ser Lys Thr Val Thr Val Gln Thr Gly Phe Gly Ala Asn Val Ala Leu His  
Asp Tyr Thr Gly Asn Gly Pro Asp Leu Arg Thr Asp Ala Tyr Gly Arg Val Thr Leu Thr Ile Pro Ala  
Asn Gly Tyr Val Ala Tyr Ser Val Pro Gly Ile Ser Gly Ser Phe Val Pro Val Glu Lys Thr Val Thr  
Gln Glu Phe Ala Gly Ala Ser Asp Leu Asp Ile Arg Pro Ala Asp Asn Thr Gln Phe Val Gln Val Gly  
Arg Ile Tyr Ala Lys Ala Asn Lys Pro Val Thr Ala Glu Leu Tyr Trp Asp Ala Lys Asp Trp Thr Thr  
Ser Thr Ser Ile Leu Leu Glu Val Arg Ser Ala Ser Gly Thr Leu Ile Thr Thr Lys Thr Val Thr Gln Leu  
Ser Ser Gln Gly Thr Arg Val Ser Phe Thr Pro Ser Ala Thr Gly Trp Tyr Val Phe Ser Ile Arg Ser Tyr  
Asn Thr Pro Ser Thr Asn Pro Lys Pro Ala Tyr Trp Leu Lys Val Thr Tyr Thr Ala Pro Gln Leu Leu  
Gln

SEQ ID NO: 131

atccgcagctttaccattgccgcgcgtggcggcgcgccggccggcagggcctggccgccttgacgctggccaccacggccctgggc  
atctcgacggcccaggcccagatgcaccgcgcacggccttcgtcatctgttcgaatggaagtggaccgacatcgcgcgcgagtgcgaga  
ccttctcgggcccgaagggttcgaggcggtgcaggtgtcggccgaacgagcacaactgggtgaccagcggtgatgtgcaccttatccg  
tggtggatgcgctaccagccggtgagctacagcctggaccgcagccgcagcgccgagttccaggacatggtcaaccgatgc  
aatgccgtgggcgtggcgtacgtggacgccgtgatcaatcacatgtccggcgccacggcgccacctcgagcgctggcgcgagctgg

Figure 16 (cont.)

agctatcacaactaccctgggctctatggcccaacgacttccaccagccggtgtgcagcatcaccaactacggggatgcgaacaatgtgcag  
cgttgagagctctcgggcttgaggacctggacactgggagcgcttatgtgcgcggcaagatgccgactatctggtgatctgtcaacatg  
ggggtaagggtctccgggtgatgcggccaaacacatcagccgaccgacctgggcgccatcatcgatcggtcaacagccgcaccggc  
gcgaaccgcccttctgtttctggaggtgattggcgcggccggcgaggcagtgagccgaaccagtacttctcgtcggcggcggccaggt  
caccgtgaccgagttcaactatgggaagcaaatcttcggcaagttcgccggtggcggcgtctggccgagctgcgcagcttcggtgaaacctg  
gggcctgatgccagcagcaaaagcgattgtttcatcgacaaccacgacaagcagcgcggtcatggcggcggtggcaactatctgacctacc  
accatggctcgacctacgatctggccaacatcttcatgtggttggccttatggctaccggcgctgatgtccagctatgcttcaaccgcagc  
acggcctacgacacgagctttggcccgccacacgacagtggtggcgccaccctggccctgggatggtggcgccagccagcggcctgc  
ttcaaccagagcatcggtggctgggtgtgtgagcaccgctggcggggcatcgcaaatatggtggccttcgcgaacgccacgctgccaactg  
gaccgtgaccgactggtgggacaacggcaacaaccagatcgcttctggcggggtgacaagggttcgtggtgatcaaccgcgaagacgc  
cgcgctgacgcgcaacttcaagaccagcctgccagccggccagctactgcgatgtcatctccggggacttcaacaatggtcagtcacggggc  
atgtggtgacggctgatgccggcggtctacgtgacgtgacggcgggcccaatggtgcggcgccatccacgtgggcggccgctgtgacg  
gcgctctcagccgccgacgaccgctcggtgacgttcaacgcgtcgccgatacctttggggacagaacctgttcgtcgtgggaaccaca  
gcgcactgggcaactggtgcggcgccggccagggcgatgacttggatttcgggtcgggcacgcgcgggaactggcgcgcggtgctca  
atttgcggccaataccacctaccaatacaagttcatcaagaaggacggggctggaaacgtgtttgggagggcggtggcaatcgctcgtga  
ccacgcgctctggggcggtatcggtgagcacggcgcccaattggcagtag

SEQ ID NO: 132

Met Pro Gln Leu Tyr Pro Leu Pro Pro Arg Trp Arg Arg Ala Ala Arg Gln Gly Leu Ala Ala Leu Thr  
Leu Ala Thr Thr Ala Leu Gly Ile Ser Thr Ala Gln Ala Gln Ser Ala Pro Arg Thr Ala Phe Val His  
Leu Phe Glu Trp Lys Trp Thr Asp Ile Ala Arg Glu Cys Glu Thr Phe Leu Gly Pro Lys Gly Phe Ala  
Ala Val Gln Val Ser Pro Pro Asn Glu His Asn Trp Val Thr Ser Gly Asp Gly Ala Pro Tyr Pro Trp  
Trp Met Arg Tyr Gln Pro Val Ser Tyr Ser Leu Asp Arg Ser Arg Ser Gly Thr Arg Ala Glu Phe Gln  
Asp Met Val Asn Arg Cys Asn Ala Val Gly Val Gly Ile Tyr Val Asp Ala Val Ile Asn His Met Ser  
Gly Gly Thr Gly Gly Thr Ser Ser Ala Gly Arg Ser Trp Ser Tyr His Asn Tyr Pro Gly Leu Tyr Gly  
Pro Asn Asp Phe His Gln Pro Val Cys Ser Ile Thr Asn Tyr Gly Asp Ala Asn Asn Val Gln Arg Cys  
Glu Leu Ser Gly Leu Gln Asp Leu Asp Thr Gly Ser Ala Tyr Val Arg Gly Lys Ile Ala Asp Tyr Leu  
Val Asp Leu Val Asn Met Gly Val Lys Gly Phe Arg Val Asp Ala Ala Lys His Ile Ser Pro Thr Asp  
Leu Gly Ala Ile Ile Asp Ala Val Asn Ser Arg Thr Gly Ala Asn Arg Pro Phe Trp Phe Leu Glu Val  
Ile Gly Ala Ala Gly Glu Ala Val Gln Pro Asn Gln Tyr Phe Ser Leu Gly Gly Gly Gln Val Thr Val  
Thr Glu Phe Asn Tyr Gly Lys Gln Ile Phe Gly Lys Phe Ala Gly Gly Gly Arg Leu Ala Glu Leu Arg  
Ser Phe Gly Glu Thr Trp Gly Leu Met Pro Ser Ser Lys Ala Ile Ala Phe Ile Asp Asn His Asp Lys  
Gln Arg Gly His Gly Gly Gly Gly Asn Tyr Leu Thr Tyr His His Gly Ser Thr Tyr Asp Leu Ala Asn  
Ile Phe Met Leu Ala Trp Pro Tyr Gly Tyr Pro Ala Leu Met Ser Ser Tyr Ala Phe Asn Arg Ser Thr  
Ala Tyr Asp Thr Ser Phe Gly Pro Pro His Asp Ser Gly Gly Ala Thr Arg Gly Pro Trp Asp Gly Gly  
Gly Ser Gln Pro Ala Cys Phe Asn Gln Ser Ile Gly Gly Trp Val Cys Glu His Arg Trp Arg Gly Ile  
Ala Asn Met Val Ala Phe Arg Asn Ala Thr Leu Pro Asn Trp Thr Val Thr Asp Trp Trp Asp Asn  
Gly Asn Asn Gln Ile Ala Phe Gly Arg Gly Asp Lys Gly Phe Val Val Ile Asn Arg Glu Asp Ala Ala  
Leu Thr Arg Asn Phe Lys Thr Ser Leu Pro Ala Gly Gln Tyr Cys Asp Val Ile Ser Gly Asp Phe Asn  
Asn Gly Gln Cys Thr Gly His Val Val Thr Val Asp Ala Gly Gly Tyr Val Thr Leu Thr Ala Gly Pro  
Asn Gly Ala Ala Ala Ile His Val Gly Ala Arg Leu Asp Gly Ala Ser Gln Pro Pro Thr Thr Ala Ser  
Val Thr Phe Asn Ala Ser Ala Asp Thr Phe Trp Gly Gln Asn Leu Phe Val Val Gly Asn His Ser Ala  
Leu Gly Asn Trp Ser Pro Ala Ala Ala Arg Pro Met Thr Trp Ile Ser Gly Ser Gly Thr Arg Gly Asn  
Trp Arg Ala Val Leu Asn Leu Pro Ala Asn Thr Thr Tyr Gln Tyr Lys Phe Ile Lys Lys Asp Gly Ala  
Gly Asn Val Val Trp Glu Gly Gly Gly Asn Arg Val Val Thr Thr Pro Ser Gly Gly Gly Ser Val Ser  
Thr Gly Gly Asn Trp Gln

SEQ ID NO: 133

atgaataatgtgaaaaagtagtgggtgtattattctataattgctaccttagttatttcttttcacacctttttcaacagcacaagctaatactgcacctg  
tcaacggaacaatgatgcaatatttcgaatgggaattaccgaatgatgggacgctttggacgaaagtaaaaaatgaagctaccaatctttctcgt

Figure 16 (cont.)

aggtattacagcgttatggctccctccagcatataaaggaacgagccaaagcgatgtcggatatggcgtgtacgatttatgacctggggaatt  
taatcaaaaaggacgatccgaacgaaatacggacaaaagcacaatatattcaagccatccaagctgccaaagccgcagggatgcaagtat  
atgcagatgttgatttaatacgaagcgggggctgacggcacagaattgtcgtacgattgaggtaaaccccttctaatacgaatacaagaacat  
ctggcacatatcaaatcaagcatggacaaaattgatttctgtgctggaaacacatactccagcttcaaatggcgtggtatcattttgacggt  
accgattgggatgaaagtcgtaaattaaatcgtatttacaattccgcggtacaggaaaagcgtgggactgggaagtcgatacagaaaacgga  
aactatgatttattaatgttcgctgatttagatggatcacctgaagttgtacagagttaaaaaactggggaataatggatgtaatacgcacaaa  
tgtagacgggatttcgttgatgccgtaaacataataaatacagctttccctgactggctaacatatgtacgtaatacaacaggaaaaattttt  
gctgttgggggaattttgactatgacgtcaataagctgcataactacattacaaaaacaatggatcgatgtcgttatttgatgcacctttgcataa  
caacttttatatcgcttccaaatcgatggatattttgacatcgcttattttgaataatacattaatgaaagatcaacctcactcgctgaacactgt  
cgataacctgatacacaaccaggtaactttacaatcatgggtagaagcttgggttaaacgcttgcttacgcctttattttaacaagacaagag  
gggtatccttgctattttacgggtgactattacggaatcccgaatacaatattccgggattaaaaagtaaaatgatccgcttttaattgctcgtcgt  
gattatgcttatggaacacaacgtgattacattgatcatcaagacattatcgatggacacgagaaggcattgatcaaaaccgaactctggactt  
gcggctttaattaccgacggccctggcggaagtaaatggatgtatcgggtaaaaacatgctgggaaagtgtttatgatttaactggaaatcga  
agtacacagtaacgattaatgcggacgggtgggagaatttaaagtaaacggcgctccgtttcgatttgggtggtctaaaacatcaaacgtca  
catttacagtcaataacgccacaacaagaaggacaaaacgtatatgtgttgcaacattccagagctaggcaattctttg

SEQ ID NO: 134

Met Asn Asn Val Lys Lys Val Trp Leu Tyr Tyr Ser Ile Ile Ala Thr Leu Val Ile Ser Phe Phe Thr Pro  
Phe Ser Thr Ala Gln Ala Asn Thr Ala Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu Trp Asp Leu  
Pro Asn Asp Gly Thr Leu Trp Thr Lys Val Lys Asn Glu Ala Thr Asn Leu Ser Ser Leu Gly Ile Thr  
Ala Leu Trp Leu Pro Pro Ala Tyr Lys Gly Thr Ser Gln Ser Asp Val Gly Tyr Gly Val Tyr Asp Leu  
Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Ile Arg Thr Lys Tyr Gly Thr Lys Ala Gln Tyr Ile  
Gln Ala Ile Gln Ala Ala Lys Ala Ala Gly Met Gln Val Tyr Ala Asp Val Val Phe Asn His Lys Ala  
Gly Ala Asp Gly Thr Glu Phe Val Asp Ala Val Glu Val Asn Pro Ser Asn Arg Asn Gln Glu Thr Ser  
Gly Thr Tyr Gln Ile Gln Ala Trp Thr Lys Phe Asp Phe Pro Gly Arg Gly Asn Thr Tyr Ser Ser Phe  
Lys Trp Arg Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Gly Lys Leu Asn Arg Ile Tyr Lys  
Phe Arg Gly Thr Gly Lys Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn Tyr Asp Tyr Leu  
Met Phe Ala Asp Leu Asp Met Asp His Pro Glu Val Val Thr Glu Leu Lys Asn Trp Gly Lys Trp  
Tyr Val Asn Thr Thr Asn Val Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Tyr Ser Phe Phe  
Pro Asp Trp Leu Thr Tyr Val Arg Asn Gln Thr Gly Lys Asn Leu Phe Ala Val Gly Glu Phe Trp Ser  
Tyr Asp Val Asn Lys Leu His Asn Tyr Ile Thr Lys Thr Asn Gly Ser Met Ser Leu Phe Asp Ala Pro  
Leu His Asn Asn Phe Tyr Ile Ala Ser Lys Ser Ser Gly Tyr Phe Asp Met Arg Tyr Leu Leu Asn Asn  
Thr Leu Met Lys Asp Gln Pro Ser Leu Ala Val Thr Leu Val Asp Asn His Asp Thr Gln Pro Gly Gln  
Ser Leu Gln Ser Trp Val Glu Ala Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Gln Glu  
Gly Tyr Pro Cys Val Phe Tyr Gly Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro Gly Leu Lys Ser  
Lys Ile Asp Pro Leu Leu Ile Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln Arg Asp Tyr Ile Asp His  
Gln Asp Ile Ile Gly Trp Thr Arg Glu Gly Ile Asp Ala Lys Pro Asn Ser Gly Leu Ala Ala Leu Ile Thr  
Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Lys His Ala Gly Lys Val Phe Tyr Asp Leu  
Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn Ala Asp Gly Trp Gly Glu Phe Lys Val Asn Gly Gly  
Ser Val Ser Ile Trp Val Ala Lys Thr Ser Asn Val Thr Phe Thr Val Asn Asn Ala Thr Thr Thr Ser  
Gly Gln Asn Val Tyr Val Val Gly Asn Ile Pro Glu Leu Gly Asn Ser Leu

SEQ ID NO: 135

gtgacaggcaccctgctttatacattctccacataaaataaccatacagctttcaatttgtgaaatgtataaaaaataaaaaatagtattgtaagc  
gttaacatccgtcattataataacttcaaacgcgtttatgttttaatgcaaacgtttgcatcctcattttttaaaagaaaggatgtgtgtcatgaattatt  
tgaaaaaagtgtggtgtattacgctatcgtcgtaccttaatacttcttttaccgcccctttcaactgcacaagccaacactgcaccagtcaacg  
gaacgatgatgcaatatttcgaatgggatttaccgaatgatggcacactttggacgaaagtaaaaaacgaagcaagcagcctttctttaggtat  
tactgcgttatggttaccctgcatacaaggaacgagccaaggggatgtcgggtatggcgtgtacgatttgatgacttaggagaatttaataca  
aaaagggacgattcgaacgaaatacggacaaaaacgcaatatttacaagccattcaagcgcaaaaaagcgtggcatgcaagtatacgtg  
atgtcgtatttaatacagaaggcggggcgagatagtacagaatgggttgacgcagtcgaagtgaatccttctaatacgaaccaagaacatctgg

Figure 16 (cont.)

cacatatcaaattcaagcatggacaaaatttgatttccctgaccgtgggaacacatactcaagctttaaattggcgctggtatcattttgacgggtacg  
gattgggatgaaagtcgaaaactaaatcgcatttacaaatttcgtggcacaggaaaagcatgggattgggaagtagacacagagaacggaac  
tatgactacttaattgttctgatttagatatggatcacctgaagtcgtgacagagctaaaaaactggggaacatggtacgtcaatacgacaaatg  
tcgatgggttctgcttagatgcagtaaagcatattaaatagcttttccagattggttaacatatgtgcgctcacaacacaaaaaatctgtttg  
cagtaggagaattttggagctacgatgtcaataaactgcataactacattacaaaaacaagtgaacctgtcgttatttgatgcgccacttcataa  
caacttttacactgcttcaaaatctagcgggtattttgacatgcgctattttgtaataatacgttgatgaaagaccagccttctcttgcggtcacactc  
gttgataatcatgacacgcaaccgggacaatctttacaatcatgggtagagccttggttaagccgcttgcttatgcctttattttgacaagacaaga  
aggatatccttgcgtattttacggcgactattacggcatccctaaatacaatattccgggattgaaaagtaaaatcgatccgcttctcattgcccgtg  
gagactacgcatacggacacacaacgtgattatattgacatcaagacattattggatggacacgggaagggaattgactcaaaaccgaactctgg  
actgcggtttaaattactgacggctcctggtggaagttaaattggatgtatgtagtataaaagcatgctggaaaagtgttttacgatctcactggaaat  
cgaagcgatacggtaacgattaatgcagacggctggggagagtttaaagtaaacggtggctccgtttccatttgggttgccaaaacatcacaaag  
tcacgtttaccgtcaacaatgcgacaacgacaagcggacaaaatgtgtatgtcgttgcaacattccagagctcggaaattggaacacagcaaa  
cgcaatcaaaatgaccccatcttcttatccaacgtggaaaacaaccattgctctccacaaggaaaagcaattggcggcggtacgccatggccctt  
ga

SEQ ID NO: 136

Val Thr Gly Thr Pro Ser Leu Tyr Ile Pro Pro His Lys Ile Thr Ile Gln Leu Ser Asn Leu Leu Lys Cys  
Ile Lys Ile Lys Asn Ser Ile Val Ser Val Asn Ile Arg His Tyr Asn Asn Phe Lys Arg Val Tyr Val Leu  
Met Gln Thr Phe Ala Ser Ser Phe Tyr Leu Lys Lys Gly Cys Val Cys Met Asn Tyr Leu Lys Lys Val  
Trp Leu Tyr Tyr Ala Ile Val Ala Thr Leu Ile Ile Ser Phe Leu Thr Pro Phe Ser Thr Ala Gln Ala Asn  
Thr Ala Pro Val Asn Gly Thr Met Met Gln Tyr Phe Glu Trp Asp Leu Pro Asn Asp Gly Thr Leu  
Trp Thr Lys Val Lys Asn Glu Ala Ser Ser Leu Ser Ser Leu Gly Ile Thr Ala Leu Trp Leu Pro Pro  
Ala Tyr Lys Gly Thr Ser Gln Gly Asp Val Gly Tyr Gly Val Tyr Asp Leu Tyr Asp Leu Gly Glu Phe  
Asn Gln Lys Gly Thr Ile Arg Thr Lys Tyr Gly Thr Lys Thr Gln Tyr Leu Gln Ala Ile Gln Ala Ala  
Lys Ser Ala Gly Met Gln Val Tyr Ala Asp Val Val Phe Asn His Lys Ala Gly Ala Asp Ser Thr Glu  
Trp Val Asp Ala Val Glu Val Asn Pro Ser Asn Arg Asn Gln Glu Thr Ser Gly Thr Tyr Gln Ile Gln  
Ala Trp Thr Lys Phe Asp Phe Pro Asp Arg Gly Asn Thr Tyr Ser Ser Phe Lys Trp Arg Trp Tyr His  
Phe Asp Gly Thr Asp Trp Asp Glu Ser Arg Lys Leu Asn Arg Ile Tyr Lys Phe Arg Gly Thr Gly Lys  
Ala Trp Asp Trp Glu Val Asp Thr Glu Asn Gly Asn Tyr Asp Tyr Leu Met Phe Ala Asp Leu Asp  
Met Asp His Pro Glu Val Val Thr Glu Leu Lys Asn Trp Gly Thr Trp Tyr Val Asn Thr Thr Asn Val  
Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Lys Tyr Ser Phe Phe Pro Asp Trp Leu Thr Tyr Val  
Arg Ser Gln Thr Gln Lys Asn Leu Phe Ala Val Gly Glu Phe Trp Ser Tyr Asp Val Asn Lys Leu His  
Asn Tyr Ile Thr Lys Thr Ser Gly Thr Met Ser Leu Phe Asp Ala Pro Leu His Asn Asn Phe Tyr Thr  
Ala Ser Lys Ser Ser Gly Tyr Phe Asp Met Arg Tyr Leu Leu Asn Asn Thr Leu Met Lys Asp Gln  
Pro Ser Leu Ala Val Thr Leu Val Asp Asn His Asp Thr Gln Pro Gly Gln Ser Leu Gln Ser Trp Val  
Glu Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe  
Tyr Gly Asp Tyr Tyr Gly Ile Pro Lys Tyr Asn Ile Pro Gly Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile  
Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln Arg Asp Tyr Ile Asp His Gln Asp Ile Ile Gly Trp Thr  
Arg Glu Gly Ile Asp Ser Lys Pro Asn Ser Gly Leu Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser  
Lys Trp Met Tyr Val Gly Lys Lys His Ala Gly Lys Val Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp  
Thr Val Thr Ile Asn Ala Asp Gly Trp Gly Glu Phe Lys Val Asn Gly Gly Ser Val Ser Ile Trp Val  
Ala Lys Thr Ser Gln Val Thr Phe Thr Val Asn Asn Ala Thr Thr Thr Ser Gly Gln Asn Val Tyr Val  
Val Gly Asn Ile Pro Glu Leu Gly Asn Trp Asn Thr Ala Asn Ala Ile Lys Met Thr Pro Ser Ser Tyr  
Pro Thr Trp Lys Thr Thr Ile Ala Leu Pro Gln Gly Lys Ala Ile Gly Gly Val Arg His Gly Pro

SEQ ID NO: 137

gtgggacgggcaggcttggcgcatcactcgaacacttccgccaaggggacatacgggtcacctctcgaactgcgtccggatcgccccgccgt  
ggccggggcggtcgagcttgaagatgtccagcggggagccgccgaggatcacccggcggtactcgccaggggcggggctcag  
cttgaagccgtggccggagccgctccaggagccagacgttgaggcccggtggtcaggaggaggtggccgtcggggctgt  
tctcgtactggcagacgcggtctcgaccagcggcgctcttcaggccgggaaccggcgggccacctcgccccggcgcttcagca

Figure 16 (cont.)

ggggccggggtgatcgtccgctgccccggcggtgggatcgtgggctgccccgggtgtcgtccgccacctgaagccgcgggtgctcgttgc  
ggggatgccgtagtagatccgctgccgagatcgaaccagaccggacagccgccctcctggaagcgcgggtgccccggcggtgctgccga  
agaagaacacctcctggcggtgttgccgaggaaccgctaccgatcacgtccgggaacagccggccagccaggaccgcaggcgaa  
acgtagaggtcggccgcgagagtggagcgtccgaaaggtgaagccgctccaaggccccgggaccatggcggcctgcccgtactcccc  
gcccctgcccctggaacagctccaccacggctccggcagggcgccggggcgaacagggcgccggttctctctgtaccagatcgtcggac  
gccgtcgaaatcgacctgggggaagcggtccgggctccccctgagacagctcggcgaccggcagccccgctctccagaaaaggaa  
gggagtcgaggacgtatgtcgtcctcggcgacatccagaggaccccggtctttgtacagccgtaaccggactggactcggcgctcc  
gccagagctcgaggagcgggcgacccactccacgtacagacggcggtcgggtccgtaggcgccgaggatgacgcgtcgcaccggag  
ctggagcgggagtgccccggacccccagcggtccaggagggtcaccgggtccggcgaggagatgcaggggcggtccagccggcg  
aaggcgccggcgccgacgaggggatggggatgggaggcatggcgggcgtaagggtatcgagcccgatcctcgtggcatccat  
ctccgaccggagatcctggaataatcgaagaaggagatcgacatgcaatcgaaacgtga

SEQ ID NO: 138

Val Gly Arg Ala Gly Leu Ala His His Ser Asn Thr Ser Ala Lys Gly Thr Tyr Gly Ser Pro Leu Glu  
Leu Arg Pro Asp Arg Pro Ala Val Ala Gly Ala Val Glu Leu Glu Asp Val Gln Arg Gly Ala Ala Ala  
Glu Asp His Pro Gly Gly Val Leu Ala Gln Gly Gly Ala Gln Leu Glu Ala Val Ala Gly Ala Ala Ser  
Gln Glu Pro Asp Val Gly Gly Pro Arg Met Ala Val Glu Glu Glu Val Ala Val Gly Ala Val Leu Val  
Leu Ala Asp Ala Gly Leu Asp Gln Arg Arg Val Leu Gln Gly Arg Glu Pro Ala Gly His Leu Gly  
Pro Gly Arg Phe Gln Gln Gly Arg Gly Asp Arg Pro Leu Ala Arg Arg Gly Ile Asp Gly Leu Ala Pro  
Gly Val Val Arg His Leu Glu Ala Ala Val Leu Val Ala Gly Asp Ala Val Val Asp Pro Leu Ala Glu  
Ile Asp Pro Asp Arg Thr Ala Ala Leu Leu Glu Ala Arg Val Ala Arg Arg Arg Ala Glu Glu Glu His  
Leu Leu Ala Gly Val Ala Glu Glu Pro Leu Thr Asp His Val Arg Glu Gln Pro Gly Gln Pro Gly Thr  
Ala Gly Glu Asp Val Glu Val Gly Arg Glu Ser Gly Ala Val Arg Lys Val Lys Pro Leu Gln Gly Pro  
Arg Asp His Gly Gly Leu Pro Val Leu Pro Ala Leu Ala Leu Glu Gln Leu His His Gly Pro Ala Gly  
Ala Pro Gly Glu Gln Gly Ala Gly Phe Leu Leu Val Pro Asp Arg Ala Asp Ala Val Glu Ile Asp Leu  
Gly Glu Ala Ala Pro Gly Leu Pro Leu Arg Gln Leu Gly Asp Arg Gln Pro Arg Val Leu Gln Lys  
Arg Lys Gly Val Ala Asp Val Ala Val Val Leu Ala Ala His Pro Glu Asp Pro Gly Pro Phe Val Gln  
Pro Val Thr Gly Leu Asp Phe Gly Val Pro Pro Glu Leu Glu Gly Ala Gly Asp Pro Leu His Val Gln  
Thr Val Gly Ser Val Gly Ala Ala Asp Asp Pro Arg Leu Ala Thr Gly Ala Gly Ala Gly Val Pro Arg  
Thr Pro Gly Val Gln Glu Gly His Pro Gly Ser Ala Ala Glu Glu Met Gln Gly Gly Pro Ala Ala Glu  
Gly Ala Gly Ala Asp Asp Gly Asp Met Gly Met Gly Gly Gly Arg Lys Val Ile Ala Ala Arg  
Ser Phe Ala Gly Ile Pro Ser Pro Thr Gly Val Ser Trp Lys Ile Arg Arg Arg Arg Ser Thr Cys Asn  
Arg Thr Glu Thr

SEQ ID NO: 139

atgaaaacattcaaccttaaacccacacttttaccttaactttgctgctgagttcgccggtattggcgccacaaaatggaactatgatgcagtatttc  
cattggtatgtgccaaatgacggcgactctggacacaagttgaaaacaatgcgccagcactatccgacaacgggtttacagcgctgtggttc  
caccagcatataaaggcgaggtgtagcaacgacgttggttacggtgtttacgatatgtatgacttaggggagtttgatcaaaaaggatcggt  
cgaactaagtacggcaccaaagaccaatatctaaatgccatcaaaagcagcacacaaaaacaatatccaaatttatggtgacgtagtgttcaacca  
tcgtggcggtgcagatggcaagtcgtgggtcgataccaagcgtgtggattggaataaccgcaatattgaacttggcgataaatggattgaagca  
tgggttgaatttagcttccaggacgtaacgataaatactcagacttccattggacgtggtatcatttgatggcgctgattgggatgacgcaggt  
aagagaaagcgatctttaaattcaaaaggtgatggtaaagcattgggtggaagtcagtttgaaaaaggcaactatgactacctatgtacgca  
gacttagacatggatcaccagaagtgaagcaagagctgaaagattgggtgaatggtacttaaacatgacgggtgttgatggcttccgaatgg  
atgcagtgaacacatcaaatatcagtagctacaagagtggtatgacttgcgtaagaaaacgggcaaagagctcttaccgttggtgagtac  
tggaactacgacgtgaacaatctgcacaactttatgactaagacttctggcagcatgtcattgtttgatgcgcctttacatatgaactctataacgt  
tcacgctctggtggaactttgatatgcgccgaatcatggtgacacttgatgaaagacaaccagtgaagcagtaaacactggttgagaacc  
atgatacgcaaccactacaggcccttagagtcctccggtgattggtggttcaaacacttgcgtacgcttcatcttctcgtgaggaaggttatcc  
gtcagctcttacgcagattactacggtgcgaatacagcgataaaggcgacgatatcaaatggtgaaagtgccttaccattgagcaattggtga  
aagcgcgtaaagattatgcttatggttaaacaacattctaccttgaccactgggatgtgattggttgacacgagaaggggatgcggaacatccg

Figure 16 (cont.)

aactctatggcgggttatcatgagtgatggctcctggcgggaacaaagtggatgtacacaggttcaccgagcacacggttatgtcgataaactaggtatt  
cgtaccgaagaagtatggactaacgctagtggatgggccgaattcccagtgaaacggcgatcggtttctgtttgggtggcgtaaataa

SEQ ID NO: 140

Met Lys Thr Phe Asn Leu Lys Pro Thr Leu Leu Pro Leu Thr Leu Leu Ser Ser Pro Val Leu Ala  
Ala Gln Asn Gly Thr Met Met Gln Tyr Phe His Trp Tyr Val Pro Asn Asp Gly Ala Leu Trp Thr Gln  
Val Glu Asn Asn Ala Pro Ala Leu Ser Asp Asn Gly Phe Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys  
Gly Ala Gly Gly Ser Asn Asp Val Gly Tyr Gly Val Tyr Asp Met Tyr Asp Leu Gly Glu Phe Asp  
Gln Lys Gly Ser Val Arg Thr Lys Tyr Gly Thr Lys Asp Gln Tyr Leu Asn Ala Ile Lys Ala Ala His  
Lys Asn Asn Ile Gln Ile Tyr Gly Asp Val Val Phe Asn His Arg Gly Gly Ala Asp Gly Lys Ser Trp  
Val Asp Thr Lys Arg Val Asp Trp Asn Asn Arg Asn Ile Glu Leu Gly Asp Lys Trp Ile Glu Ala Trp  
Val Glu Phe Ser Phe Pro Gly Arg Asn Asp Lys Tyr Ser Asp Phe His Trp Thr Trp Tyr His Phe Asp  
Gly Val Asp Trp Asp Asp Ala Gly Lys Glu Lys Ala Ile Phe Lys Phe Lys Gly Asp Gly Lys Ala Trp  
Asp Trp Glu Val Ser Ser Glu Lys Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp  
His Pro Glu Val Lys Gln Glu Leu Lys Asp Trp Gly Glu Trp Tyr Leu Asn Met Thr Gly Val Asp Gly  
Phe Arg Met Asp Ala Val Lys His Ile Lys Tyr Gln Tyr Leu Gln Glu Trp Ile Asp Tyr Leu Arg Lys  
Lys Thr Gly Lys Glu Leu Phe Thr Val Gly Glu Tyr Trp Asn Tyr Asp Val Asn Asn Leu His Asn  
Phe Met Thr Lys Thr Ser Gly Ser Met Ser Leu Phe Asp Ala Pro Leu His Met Asn Phe Tyr Asn Ala  
Ser Arg Ser Gly Gly Asn Phe Asp Met Arg Arg Ile Met Asp Gly Thr Leu Met Lys Asp Asn Pro  
Val Lys Ala Val Thr Leu Val Glu Asn His Asp Thr Gln Pro Leu Gln Ala Leu Glu Ser Pro Val Asp  
Trp Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Leu Arg Glu Glu Gly Tyr Pro Ser Val Phe Tyr  
Ala Asp Tyr Tyr Gly Ala Gln Tyr Ser Asp Lys Gly His Asp Ile Asn Met Val Lys Val Pro Tyr Ile  
Glu Gln Leu Val Lys Ala Arg Lys Asp Tyr Ala Tyr Gly Lys Gln His Ser Tyr Leu Asp His Trp Asp  
Val Ile Gly Trp Thr Arg Glu Gly Asp Ala Glu His Pro Asn Ser Met Ala Val Ile Met Ser Asp Gly  
Pro Gly Gly Thr Lys Trp Met Tyr Thr Gly Ser Pro Ser Thr Arg Tyr Val Asp Lys Leu Gly Ile Arg  
Thr Glu Glu Val Trp Thr Asn Ala Ser Gly Trp Ala Glu Phe Pro Val Asn Gly Gly Ser Val Ser Val  
Trp Val Gly Val Lys

SEQ ID NO: 141

atgaaccaataaataccctactcatatccgcccttgcgtgtttgtttcagttccgcgacttacgccgatactattttgcacgcgttcaattggaagt  
attcagatgtgacggccaacgcgaatcaaattgctcaagctggttataagaaagtgcgtgttgcgcctgcaatgaaatcgagtgccagccaatgg  
tgggctcgctatcaacctcaagatctacgcactatcgattctcctttgggcaataaacaagatttagccgcaatgattgccgcactcaaaggtgtg  
ggcgtcgatgtgatgccgatgtgttactcaaccatatggcgaatgaaagctggaagcgaagtgaattacacctggcacagaagtgctaa  
acgattatgctagccgttcaagctactatgctgaccagactctgtttggcaacctagcacaaggttatgtgtcagcgaacgactttcatccagcgg  
gctgtatttcagattggaacgaccctggtcatgttcagattggcgtttgtgtggcgagatggtgatgtaggtttacctgaccttgatccaaacaac  
tgggtggtttcacaacagcgtttgtatctgaaagcgctaaaagatatgggcatcaaagggttccgaattgatgcagtgaagcatgagccaata  
ccaaatcgatcaggtattcacgtctgaaattactgcgaacatgcatgttttggtgaagtgattactagcgggtggagcaggggaatagcggctatg  
aatcgttcttagcgccttacctgaataataactactctgcctacgattcccgtgtttgcacgattcgctcggcattttctatggggggcggttt  
aatcaactgcatgatcctaaagcgtagggtcaggcacttgatgataatcgctcgatcacccttgcgacacatgatattccaaccaatgacgg  
cttccgctaccaaattatggaccacacaagacgagcagcttgcttacgcgtatatccttggttaaagacgggtggcacgccgctgatctacagtgatg  
atcttctgattctgaagacaaggataaacggtcgttggggcaatgtttggaacagttcgacaatgaaaacatgttgagcttcataacgcgatgc  
aaggcaaaacaatgacgatgattctagcgaccattgcactttgtttgaagcgtggcaagaaggtgttggtggttataacaagtgtgtgtaa  
cgcggtggcgtgacggttgatacctaccaacatgagtttaattggcatgttcaatacaagacgtgtaagcagcgcaacagaaaccgtgacttct  
cgttaccatacgttcaatctaccaccagcagtcgcggtatgtttaagctgatg

SEQ ID NO: 142

Met Lys Pro Ile Asn Thr Leu Leu Ile Ser Ala Leu Ala Val Cys Ser Phe Ser Ser Ala Thr Tyr Ala  
Asp Thr Ile Leu His Ala Phe Asn Trp Lys Tyr Ser Asp Val Thr Ala Asn Ala Asn Gln Ile Ala Gln  
Ala Gly Tyr Lys Lys Val Leu Val Ala Pro Ala Met Lys Ser Ser Gly Ser Gln Trp Trp Ala Arg Tyr  
Gln Pro Gln Asp Leu Arg Thr Ile Asp Ser Pro Leu Gly Asn Lys Gln Asp Leu Ala Ala Met Ile Ala

Parameter	Value	Unit	Parameter	Value	Unit
$\alpha$	0.001		$\beta$	0.001	
$\gamma$	0.001		$\delta$	0.001	
$\epsilon$	0.001		$\zeta$	0.001	
$\eta$	0.001		$\theta$	0.001	
$\iota$	0.001		$\kappa$	0.001	
$\lambda$	0.001		$\mu$	0.001	
$\nu$	0.001		$\xi$	0.001	
$\omicron$	0.001		$\pi$	0.001	
$\rho$	0.001		$\sigma$	0.001	
$\tau$	0.001		$\upsilon$	0.001	
$\phi$	0.001		$\chi$	0.001	
$\psi$	0.001		$\omega$	0.001	
$\Omega$	0.001		$\Theta$	0.001	
$\Phi$	0.001		$\Psi$	0.001	
$\Upsilon$	0.001		$\Xi$	0.001	
$\Lambda$	0.001		$\Sigma$	0.001	
$\Pi$	0.001		$\Upsilon$	0.001	
$\Gamma$	0.001		$\Delta$	0.001	
$\Sigma$	0.001		$\Omega$	0.001	
$\Delta$	0.001		$\Theta$	0.001	
$\Omega$	0.001		$\Phi$	0.001	
$\Theta$	0.001		$\Psi$	0.001	
$\Phi$	0.001		$\Xi$	0.001	
$\Psi$	0.001		$\Sigma$	0.001	
$\Xi$	0.001		$\Upsilon$	0.001	
$\Sigma$	0.001		$\Delta$	0.001	
$\Upsilon$	0.001		$\Omega$	0.001	
$\Delta$	0.001		$\Theta$	0.001	
$\Omega$	0.001		$\Phi$	0.001	
$\Theta$	0.001		$\Psi$	0.001	
$\Phi$	0.001		$\Xi$	0.001	
$\Psi$	0.001		$\Sigma$	0.001	
$\Xi$	0.001		$\Upsilon$	0.001	
$\Sigma$	0.001		$\Delta$	0.001	
$\Upsilon$	0.001		$\Omega$	0.001	
$\Delta$	0.001		$\Theta$	0.001	
$\Omega$	0.001		$\Phi$	0.001	
$\Theta$	0.001		$\Psi$	0.001	
$\Phi$	0.001		$\Xi$	0.001	
$\Psi$	0.001		$\Sigma$	0.001	
$\Xi$	0.001		$\Upsilon$	0.001	
$\Sigma$	0.001		$\Delta$	0.001	
$\Upsilon$	0.001		$\Omega$	0.001	
$\Delta$	0.001		$\Theta$	0.001	
$\Omega$	0.001		$\Phi$	0.001	
$\Theta$	0.001		$\Psi$	0.001	
$\Phi$	0.001		$\Xi$	0.001	
$\Psi$	0.001		$\Sigma$	0.001	
$\Xi$	0.001		$\Upsilon$	0.001	
$\Sigma$	0.001		$\Delta$	0.001	
$\Upsilon$	0.001		$\Omega$	0.001	
$\Delta$	0.001		$\Theta$	0.001	
$\Omega$	0.001		$\Phi$	0.001	
$\Theta$	0.001		$\Psi$	0.001	
$\Phi$	0.001		$\Xi$	0.001	
$\Psi$	0.001		$\Sigma$	0.001	
$\Xi$	0.001		$\Upsilon$	0.001	
$\Sigma$	0.001		$\Delta$	0.001	
$\Upsilon$	0.001		$\Omega$	0.001	
$\Delta$	0.001		$\Theta$	0.001	
$\Omega$	0.001		$\Phi$	0.001	
$\Theta$	0.001		$\Psi$	0.001	
$\Phi$	0.001		$\Xi$	0.001	
$\Psi$	0.001		$\Sigma$	0.001	
$\Xi$	0.001		$\Upsilon$	0.001	
$\Sigma$	0.001		$\Delta$	0.001</	

SEQ ID NO: 143

SEQ ID NO: 144

SEQ ID NO: 144

Met Pro Lys Ser Thr Phe Thr Lys Ser Ile Thr Lys Ser Leu Leu Ala Thr Ser Val Val Val Ser Leu  
Leu Pro Ala Tyr Ala Gln Ala Asp Thr Ile Leu His Ala Phe Asn Trp Lys Tyr Ser Asp Ile Thr Arg  
Gln Ala Glu Gln Ile Ala Gln Ala Gly Tyr Lys Lys Val Leu Ile Ser Pro Pro Leu Lys Ser Thr Gly Pro  
Gln Trp Trp Ala Arg Tyr Gln Pro Gln Asp Ile Arg Val Ile Asp Ser Pro Val Gly Asn Lys Gln Asp  
Leu Gln Ala Leu Ile Ala Ala Leu Lys Ala Gln Gly Val Glu Val Tyr Ala Asp Ile Val Leu Asn His  
Met Ala Asn Glu Ser Trp Lys Arg Asp Asp Leu Asn Tyr Pro Gly Ser Asp Leu Leu Thr Gln Tyr Ser  
Gln Asn Met Ala Tyr Met Asn Gln Gln Lys Leu Phe Gly Asp Leu Glu Gln Asn Gln Phe Ser Ala  
Asn Asp Phe His Pro Ala Gly Cys Ile Thr Asp Trp Ser Asn Pro Gly His Val Gln Tyr Trp Arg Leu  
Cys Gly Gly Asn Gly Asp Thr Gly Leu Pro Asp Leu Asp Pro Asn Ser Trp Val Ile Asp Gln Gln Lys  
Arg Tyr Leu Arg Ala Leu Lys Asp Met Gly Ile Lys Gly Phe Arg Val Asp Ala Val Lys His Met Ser  
Asp Tyr Gln Ile Asn Gln Val Phe Thr Pro Asp Ile Ile Ala Gly Leu His Val Phe Gly Glu Val Ile Thr  
Ser Gly Gly Lys Gly Ser Asn Asp Tyr His Ser Phe Leu Glu Pro Tyr Leu Asn Asn Thr Asn His Ala  
Ala Tyr Asp Phe Pro Leu Phe Ala Ser Ile Arg Asn Ala Phe Ser Tyr His Gly Ser Leu Ser Gln Leu  
His Asp Pro Gln Ala Tyr Gly Gln Ala Leu Pro Asn Asp Arg Ala Ile Thr Phe Thr Ile Thr His Asp

Figure 16 (cont.)

Ile Pro Thr Asn Asp Gly Phe Arg Tyr Gln Ile Met Asp Pro Thr Ser Glu Lys Leu Ala Tyr Ala Tyr  
Ile Leu Gly Lys Asp Gly Gly Ser Pro Leu Ile Tyr Ser Asp Ala Leu Asp Pro Ser Glu Asp Lys Asp  
Lys Gly Arg Trp Arg Asp Val Trp Asn Gln Glu Tyr Met Val Asn Met Ile Ser Phe His Asn Lys Val  
Gln Gly Lys Ser Met Glu Val Met Tyr Ser Asp Gln Cys Leu Leu Val Phe Lys Arg Glu Lys Gln  
Gly Leu Val Gly Ile Asn Lys Cys Ala Glu Ser Arg Thr Tyr Thr Ile Asp Thr His Arg Phe Glu Phe  
Asn Trp Tyr Gln Pro Tyr Asn Asp Thr Leu Ser Gln His Ser Glu Thr Phe Ser Ser Arg Tyr His Ala  
Leu Thr Ile Pro Ala Gln Thr Ala Arg Met Leu Ala Leu

SEQ ID NO: 145

atgttgaagaggattacggtagtctgtttatttatttttgccttttcctaataatatatgggaggaataaggcggaagcagcaacgataaataatgga  
acattaatgcagtattttgagtggtacgctccgaatgatgggaatcattggaatcggttgcttatgatgctgaaagttagctcataagggaatcac  
atctgtatggataccacctgcataaaagggaacttcgaaaatgatgtagggtatggggcctatgatttatacgatttaggggagttcaatcaaaaa  
ggaacgggtgcggacgaaatatgggacaaaggcacagttgaaatctgcaattgacgctttacataagcaaacatcgacgtatagcggtagta  
gttatgaatcataaagggtggggctgattatactgaaaccgtaacagctgttgaggtagaccgtaacaatcgaaatattgaagtatcaggtgattatg  
aaattagtcgtggacgggtttaactttccagggcgagagatgcttattcatttcaaatggaaatgggtatcattttgacggaacggattgggat  
gaaggaaaggaaattaaaccgaattataaatttaggggtataggtaaagcgtgggactgggaagtgtctagcgaaaatggaaattatgattttg  
atgtatgcagatcttgattttgatcatccagatgttgcgaatgaaatgaaaagttggggaacgtggtatgcgaatgaattaaatttagatggattcgt  
ttagatgctgttaaacatattgatcatgaatatttacgcgattgggtaaatcatgtcagacagcaaacggggaagaaatgtttacgggtggctgaat  
attggcaaaatgataccagactttaacaattatttggcgaaagtcaattataatcaatctgtatttgatgcaccgcttcattacaattttcattatgctt  
caacaggaaatgggaattatgatagagaaatattttaaattggaacagtaatgaaaaatcatcctgcactcgcagttactctcgttgagaatcatga  
ttctaacctgggcaatcattggaatctgtagtaagtcggtgttaagccgctggcatatgcatttatttaactcgtgcagagggctatccttcagt  
ttttatgggtattactatgggacaagcggaaatagtagttatgaaattccagcggttaaaagataaaattgatccaattttgacggcacgaaaaaact  
ttgcatatggtagcagcgtgattatttagaccatccagatgtgattggctggacaagagaaggagatagtgatcatgtaagtctggttagcgg  
cattaatctccgatggaccaggagatcaaaagtggatggatgttggaaagaataacgctggggaagtatggtacgatattacgggtaatcaaac  
aaatactgtaacaattaataaagatggatcggggcaattccatgtaagtggaggctctgtttctatatatgttcaacagtaa

SEQ ID NO: 146

Met Leu Lys Arg Ile Thr Val Val Cys Leu Leu Phe Ile Leu Leu Phe Pro Asn Ile Tyr Gly Arg Asn  
Lys Ala Glu Ala Ala Thr Ile Asn Asn Gly Thr Leu Met Gln Tyr Phe Glu Trp Tyr Ala Pro Asn Asp  
Gly Asn His Trp Asn Arg Leu Arg Tyr Asp Ala Glu Ser Leu Ala His Lys Gly Ile Thr Ser Val Trp  
Ile Pro Pro Ala Tyr Lys Gly Thr Ser Gln Asn Asp Val Gly Tyr Gly Ala Tyr Asp Leu Tyr Asp Leu  
Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr Lys Ala Gln Leu Lys Ser Ala Ile  
Asp Ala Leu His Lys Gln Asn Ile Asp Val Tyr Gly Asp Val Val Met Asn His Lys Gly Gly Ala Asp  
Tyr Thr Glu Thr Val Thr Ala Val Glu Val Asp Arg Asn Asn Arg Asn Ile Glu Val Ser Gly Asp Tyr  
Glu Ile Ser Ala Trp Thr Gly Phe Asn Phe Pro Gly Arg Arg Asp Ala Tyr Ser Asn Phe Lys Trp Lys  
Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Gly Arg Lys Leu Asn Arg Ile Tyr Lys Phe Arg Gly  
Ile Gly Lys Ala Trp Asp Trp Glu Val Ser Ser Glu Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp  
Leu Asp Phe Asp His Pro Asp Val Ala Asn Glu Met Lys Ser Trp Gly Thr Trp Tyr Ala Asn Glu Leu  
Asn Leu Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Asp His Glu Tyr Leu Arg Asp Trp Val Asn  
His Val Arg Gln Gln Thr Gly Lys Glu Met Phe Thr Val Ala Glu Tyr Trp Gln Asn Asp Ile Gln Thr  
Leu Asn Asn Tyr Leu Ala Lys Val Asn Tyr Asn Gln Ser Val Phe Asp Ala Pro Leu His Tyr Asn  
Phe His Tyr Ala Ser Thr Gly Asn Gly Asn Tyr Asp Met Arg Asn Ile Leu Asn Gly Thr Val Met Lys  
Asn His Pro Ala Leu Ala Val Thr Leu Val Glu Asn His Asp Ser Gln Pro Gly Gln Ser Leu Glu Ser  
Val Val Ser Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Ala Glu Gly Tyr Pro Ser  
Val Phe Tyr Gly Asp Tyr Tyr Gly Thr Ser Gly Asn Ser Ser Tyr Glu Ile Pro Ala Leu Lys Asp Lys  
Ile Asp Pro Ile Leu Thr Ala Arg Lys Asn Phe Ala Tyr Gly Thr Gln Arg Asp Tyr Leu Asp His Pro  
Asp Val Ile Gly Trp Thr Arg Glu Gly Asp Ser Val His Ala Lys Ser Gly Leu Ala Ala Leu Ile Ser  
Asp Gly Pro Gly Gly Ser Lys Trp Met Asp Val Gly Lys Asn Asn Ala Gly Glu Val Trp Tyr Asp Ile  
Thr Gly Asn Gln Thr Asn Thr Val Thr Ile Asn Lys Asp Gly Ser Gly Gln Phe His Val Ser Gly Gly  
Ser Val Ser Ile Tyr Val Gln Gln

Figure 16 (cont.)

SEQ ID NO: 147

atgagctaaataactttaaggtaaaactgcttagtttgcgtgtcttctgctgtattgtcactggctccaaatttagccaatgctgcaaatttgaag  
tgagatgggtgataatccatccgtttcagtgacatatgacaatatagcaaaagagtgtacagagtaccttggtccagccggttgacgggtgaca  
gatttcccagccagcgggaacataagcgggctgaaggagtaggtgggcttatatcagccggttaattataagaattttacaacatgaccggta  
acgaggagcagcttaaggcaatgatcaagacctgtaatgatcaggtgttaagggtgtcgtgacgctgttttaacaaaaggctacagacgg  
tgtaggctggggcgggttaacttgaggtataagaactacctgacggattctccggatcagattccatggagactgtccattgacaaaagctat  
actgatgcaaatatgtcagaacctgtgcactctcaggtatgccggacgttgccacagataactccgctactcaggaaaagattgcagattacct  
cgcttcttaataatgaatatgggggtctatgggttccgtattgacgctgcaaaacacatgggatacaacgatatcaactccattcttcaaaaactgcac  
agaagactggaagaagacctcctgcatactggaagtaatcgagcggtaacgaagctgccgacattcagccggacaagtatactttattga  
gaatgcggtgttaactgacttcgggtatgtctgggatgcaaatgagagtttggaaagggttaattacggtaaggcactggaactcagtacctggct  
cggtgcaaatcagaacattcgtaaacaatcatgatgatgaatggggcagatgctcagccggtagctgctcaatgaaaactcagaattatgctg  
attataatctggctcagtcctggcttgctgtatggcctgtaggtacagtaagacagatatattccggtattcattccctgtaaaagataatgacctta  
tcgctcagtgatgcaactcatgatcaggcgggcctcttggtgccgaccgtgtgaagggtggctggtgtgtcagcaccgtgtgtcttctgttct  
caattcccaagatttgcgagagctaccagaggtactgctgtatcaaccaagggtattgacaatgggtgcttgggttgaacagaggaaagcaag  
gttttatgcacagaatactaccaacagctctataaccagacattctctgttgaaagtacctgcaggaaattactgtgatatcttaggaacatcagat  
cctaagagcaatccatgcggagcagacgttgtcgaagcggcggttaaggctacctttactattctgcaaaagacagctgtggctatctgtacaga  
ctcagactggtgcggcaaggggggttgatccttctgtaaaagtacccgaccgggtgctgctgtgttgaagggggaaaccaccgttaattggtgtgt  
gcgtcagctggtgtaatgcgcattcatcaaatgaggaatgcacctgtgtattgaatccgaatgatgccaactgcaggtgtatgtgaacctacca  
agggtaaacctgtttacgccggtacttcaaacgggtggaacaggatccttaacatataaccgtaaaacaggtttcttgactattaatctgactctt  
gacggtgcaggtgataccagcggagctcagcgttcaaggttacagacggatgttcatggaccggaacagttacggttcttcaggtactgccg  
gaaagttggatgtaatacatcatcaaccggcgatgaacctgtgtcttctgtgtgattatgttcttccattaacgataagaccatggaatacat  
tcaccaaggcagatgaagtaactaatcagccaccgggtgcacatttaccgcagagttacagggtctgaccgtttctttgccaataattcatccga  
ccctgagaatgatgaattaacctacagctggaatttcggtaattggtaaaacatcatccgagaaagctcttagcataacctatgaagaatccggtta  
agtatactgttactttaagggttactgattcagtaataacactgatacttactaaagataactgtaacagcaccttctagtggcaagtacttaaa  
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gtgtcgacctgcaggcgcgcgagctc

SEQ ID NO: 148

Met Ser Leu Asn Asn Phe Lys Val Lys Leu Leu Ser Phe Ala Val Ser Ser Ala Val Leu Ser Leu Ala  
Pro Asn Leu Ala Asn Ala Ala Asn Phe Glu Ser Glu Met Val Ile Ile His Pro Phe Gln Trp Thr Tyr  
Asp Asn Ile Ala Lys Glu Cys Thr Glu Tyr Leu Gly Pro Ala Gly Phe Asp Gly Val Gln Ile Ser Gln  
Pro Ala Glu His Lys Arg Ala Glu Gly Val Trp Trp Ala Val Tyr Gln Pro Val Asn Tyr Lys Asn Phe  
Thr Thr Met Thr Gly Asn Glu Glu Gln Leu Lys Ala Met Ile Lys Thr Cys Asn Asp Ala Gly Val Lys  
Val Phe Ala Asp Ala Val Phe Asn Gln Lys Ala Thr Asp Gly Val Gly Trp Gly Gly Ser Thr Trp Ser  
Tyr Lys Asn Tyr Pro Asp Gly Phe Ser Gly Ser Asp Phe His Gly Asp Cys Ser Ile Asp Lys Ser Tyr  
Thr Asp Ala Asn Asn Val Arg Thr Cys Ala Leu Ser Gly Met Pro Asp Val Tyr Gly Phe Arg Ile Asp  
Ala Ala Lys His Met Gly Tyr Asn Asp Ile Asn Ser Ile Leu Ser Lys Thr Ala Gln Lys Thr Gly Arg  
Arg Pro Pro Ala Tyr Leu Glu Val Ile Gly Ala Gly Asn Glu Ala Ala Asp Ile Gln Pro Asp Lys Tyr  
Thr Phe Ile Glu Asn Ala Val Val Thr Asp Phe Gly Tyr Val Trp Asp Ala Asn Glu Ser Phe Gly Lys  
Gly Asn Tyr Gly Lys Ala Leu Glu Leu Ser Thr Trp Leu Gly Ala Asn Ser Glu Thr Phe Val Asn Asn  
His Asp Asp Glu Trp Gly Arg Cys Ser Ala Gly Ser Cys Ser Met Lys Thr Gln Asn Tyr Ala Asp Tyr  
Asn Leu Ala Gln Ser Trp Leu Ala Val Trp Pro Val Gly Thr Val Arg Gln Ile Tyr Ser Gly Tyr Ser  
Phe Pro Val Lys Asp Asn Asp Pro Tyr Arg Val Ser Asp Ala Thr His Asp Gln Gly Gly Pro Leu Gly  
Ala Asp Arg Cys Glu Gly Gly Trp Leu Cys Gln His Arg Val Ser Phe Val Leu Asn Ser Pro Arg Phe  
Ala Arg Ala Thr Arg Gly Thr Ala Val Ser Thr Lys Gly Phe Asp Asn Gly Ala Leu Trp Phe Asn Arg  
Gly Ser Lys Gly Phe Tyr Ala Gln Asn Thr Thr Asn Ser Pro Ile Thr Gln Thr Phe Ser Val Glu Val  
Pro Asp Gly Asn Tyr Cys Asp Ile Leu Gly Thr Ser Asp Pro Lys Ser Asn Pro Cys Gly Ala Asp Val  
Val Val Ser Gly Gly Lys Ala Thr Phe Thr Ile Pro Ala Lys Thr Ala Val Ala Ile Cys Thr Asp Ser

Figure 16 (cont.)

Asp Trp Cys Gly Lys Gly Val Asp Pro Cys Glu Ser Asp Pro Thr Gly Ala Ala Cys Val Cys Lys Gly  
Glu Thr Thr Val Asn Gly Val Cys Val Ser Trp Cys Asn Ala His Ser Ser Asn Glu Glu Cys Thr Cys  
Val Leu Asn Pro Asn Asp Ala Asn Cys Gln Ala Asp Ile Glu Pro Thr Lys Gly Lys Leu Cys Tyr Ala  
Gly Thr Ser Asn Gly Trp Lys Gln Asp Pro Leu Thr Tyr Asn Arg Lys Thr Gly Phe Trp Thr Ile Asn  
Leu Thr Leu Asp Gly Ala Gly Asp Thr Ser Gly Ala Gln Arg Phe Lys Val Thr Asp Gly Cys Ser Trp  
Thr Gly Thr Val Tyr Gly Ser Ser Gly Thr Ala Gly Lys Leu Asp Val Asn Thr Ser Ser Thr Gly Asp  
Glu Pro Val Ser Leu Val Gly Asp Tyr Val Leu Ser Ile Asn Asp Lys Thr Met Glu Tyr Thr Phe Thr  
Lys Ala Asp Glu Val Thr Asn Gln Pro Pro Val Ala Ser Phe Thr Ala Thr Val Asn Gly Leu Thr Val  
Ser Phe Ala Asn Asn Ser Ser Asp Pro Glu Asn Asp Glu Leu Thr Tyr Ser Trp Asn Phe Gly Asn Gly  
Lys Thr Ser Ser Glu Lys Ala Pro Ser Ile Thr Tyr Glu Glu Ser Gly Lys Tyr Thr Val Thr Leu Lys  
Val Thr Asp Ser Ala Asn Asn Thr Asp Thr Phe Thr Lys Asp Ile Thr Val Thr Ala Pro Ser Ser Gly  
Lys Tyr Leu Lys Val Ala Val Arg Gly Ser His Asp Asn Tyr Gly Thr Asp Leu Leu Thr Lys Asn Gly  
Ser Asp Trp Thr Gly Val Phe Glu Phe Phe Gly Ser Thr Ser Val Asp Leu Gln Ala Arg Glu Leu

SEQ ID NO: 149

atgatcttaagtaattttaaggtaaacttcttagtttctgtgtcttctgtgtactgacactggctgcaaatgtcgccaatgccaagaattatgaaa  
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gatttcccaggcggtgagcataaagatgccggtggtgcatggtgggtacacctaccgctgtaactcaagagtttactaccatggttgta  
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tgtaggataggtggttcaactttcggaaattataattatctgacggattaccagtgatgatttcatcataataactgcagtataggttaataatt  
cagatgcatgggtagtaagattctgtgacctcagtggtcatgccgatatagcaactgataacgacagtaccagaaataagattgctgattactcg  
ccagccttatgaatatgggggtatagcgattccgtattgatgctgccaagcactttagctatgatgatagacgctattgtagagaaaacagcaa  
ccaaagcaggcaggagacctcctgtctatatggaagggtatcggtatccgggtcaagaggcggtatgatccagccgaacaagtatacatgga  
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cgtttgtgctcaattctctagatttgcaagagcaaccagagggacagctgttactactaaagggtttgatgacggagcctttgtggttaacagagg  
aagcaagggtctctatgccagaatactaccggcagttctataactcatacattctcagttgaattacctgatgaaattactgtgataccttgag  
caaccgatccgaagaataatccttgcggagcggtatgactgtaagcggaggttaaagcaacctttaccattccggcaagaccgccgtagcta  
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aacggcgatgtgtaagctgtgtgtaatgctcactcatctaatgaagaatgtcctgtgtgctaaatcctaatacagctgagtgtagccgacatt  
gagccgaccaagggttaaactctgctatgtaggtacctccaacaagtggactcaggaaacctttaacctataatcgcaagaccggttctggactct  
caacgttgaacttgacggttaaggggataaccagcggggcgagcgctttaaagttaccgacggctgtcatggcagggtactgtttacggttca  
tcaggagtagaaggcagacttgacgtaaatacttcagccaccggagatgaaccgggttactgacaggtaaatatgttcttccataaatgataag  
accatggaatacacattcattcctgcaggcagtggaacaagcctccggttgcgtcatttactccgactgttaaagatctgactgtatctttgtcaa  
taattcatccgacctgagaatgatgaattaacctacagctggaatttcggttaacggtaaaacctcatctgaaaagaatccgagtggttacatatgat  
aaagccggtaaatatactgtttcactcaaagtaaccgatactgcaaacacactgataccaaaacactggaatcgatttaacatcctgttaacg  
gaaaatattccaaggttgacgtcagaggttcacatgataactacggaacaaatctgtaaccagggaatggttcagaatggaccgggtatcttgaatt  
cagtaagacaacaaattcaagcttgaagctctgctcctgcagctgaccagtgatcttctcggcggtaatcgaggtgagggcattgactgcct  
ccggtggttatctctcctcgcggaagggtataactataaagttaatgaggaaagcaagggttctactgcaggcgatgttgactgcaccggg

SEQ ID NO: 150

Met Ile Leu Ser Asn Phe Lys Val Lys Leu Leu Ser Phe Ala Val Ser Ser Ala Val Leu Thr Leu Ala  
Ala Asn Val Ala Asn Ala Lys Asn Tyr Glu Ser Glu Met Val Ile Ile His Pro Phe Gln Trp Thr Tyr  
Asp Asn Ile Ala Lys Glu Cys Thr Glu Tyr Leu Gly Pro Ala Gly Phe Asp Gly Val Gln Ile Ser Gln  
Ala Ala Glu His Lys Asp Ala Gly Gly Ala Trp Trp Gly Thr Tyr Gln Pro Val Asn Phe Lys Ser Phe  
Thr Thr Met Val Gly Asn Glu Glu Gln Leu Arg Ala Met Ile Lys Thr Cys Asn Glu Ala Gly Val Lys  
Val Phe Ala Asp Ala Val Ile Asn Gln Lys Ala Gly Asp Gly Val Gly Ile Gly Gly Ser Thr Phe Gly  
Asn Tyr Asn Tyr Pro Asp Gly Phe Thr Ser Asp Asp Phe His His Asn Asn Cys Ser Ile Gly Asn Asn

Figure 16 (cont.)

Tyr Ser Asp Ala Trp Val Val Arg Phe Cys Asp Leu Ser Gly Met Pro Asp Ile Ala Thr Asp Asn Asp  
Ser Thr Arg Asn Lys Ile Ala Asp Tyr Phe Ala Ser Leu Met Asn Met Gly Val Tyr Gly Phe Arg Ile  
Asp Ala Ala Lys His Phe Ser Tyr Asp Asp Ile Asp Ala Ile Val Glu Lys Thr Ala Thr Lys Ala Gly  
Arg Arg Pro Pro Val Tyr Met Glu Val Ile Gly Asn Pro Gly Gln Glu Ala Asp Asp Ile Gln Pro Asn  
Lys Tyr Thr Trp Ile Asp Asn Ala Val Val Thr Asp Phe Thr Tyr Ala Asn Ser Met His Asn Ile Phe  
Asn Gly Ser Gly Tyr Ala Lys Ala Leu Asn Met Gly Leu Gly His Val Asp Ala Glu Asn Ala Glu Val  
Phe Ile Ser Asn His Asp Asn Glu Trp Gly Arg Lys Ser Ala Gly Ser Cys Ser Ile Arg Thr Gln Asn  
Asn Pro Asp Tyr His Leu Ala Gln Ser Trp Leu Ala Val Trp Pro Leu Gly Lys Val Arg Gln Ile Tyr  
Ser Ala Tyr Gln Phe Pro Val Phe Glu Asp Ser Cys Glu Arg Val Ser Gln Gln Ala His Asp Gln Gly  
Gly Pro Ile Gly Ala Ala Arg Cys Glu Gly Gly Trp Leu Cys Gln His Arg Val Pro Phe Val Leu Asn  
Ser Pro Arg Phe Ala Arg Ala Thr Arg Gly Thr Val Val Thr Thr Lys Gly Phe Asp Asp Gly Ala Leu  
Trp Phe Asn Arg Gly Ser Lys Gly Phe Tyr Ala Gln Asn Thr Thr Gly Ser Ser Ile Thr His Thr Phe  
Ser Val Glu Leu Pro Asp Gly Asn Tyr Cys Asp Ile Leu Gly Ala Thr Asp Pro Lys Asn Asn Pro Cys  
Gly Ala Asp Val Thr Val Ser Gly Gly Lys Ala Thr Phe Thr Ile Pro Ala Lys Thr Ala Val Ala Ile  
Cys Thr Asp Glu Lys Trp Cys Gly Lys Gly Val Asp Pro Cys Glu Ser Asp Pro Thr Gly Ser Ala Cys  
Val Cys Lys Gly Glu Thr Thr Val Asn Gly Val Cys Val Ser Trp Cys Asn Ala His Ser Ser Asn Glu  
Glu Cys Ala Cys Val Leu Asn Pro Asn Asp Ala Glu Cys Gln Ala Asp Ile Glu Pro Thr Lys Gly Lys  
Leu Cys Tyr Val Gly Thr Ser Asn Lys Trp Thr Gln Glu Pro Leu Thr Tyr Asn Arg Lys Thr Gly Phe  
Trp Thr Leu Asn Val Glu Leu Asp Gly Lys Gly Asp Thr Ser Gly Ala Gln Arg Phe Lys Val Thr  
Asp Gly Cys Ser Trp Gln Gly Thr Val Tyr Gly Ser Ser Gly Val Glu Gly Arg Leu Asp Val Asn Thr  
Ser Ala Thr Gly Asp Glu Pro Val Ser Leu Thr Gly Lys Tyr Val Leu Ser Ile Asn Asp Lys Thr Met  
Glu Tyr Thr Phe Ile Pro Ala Gly Ser Gly Asn Lys Pro Pro Val Ala Ser Phe Thr Pro Thr Val Lys  
Asp Leu Thr Val Ser Phe Val Asn Asn Ser Ser Asp Pro Glu Asn Asp Glu Leu Thr Tyr Ser Trp Asn  
Phe Gly Asn Gly Lys Thr Ser Ser Glu Lys Asn Pro Ser Val Thr Tyr Asp Lys Ala Gly Lys Tyr Thr  
Val Ser Leu Lys Val Thr Asp Thr Ala Asn Asn Thr Asp Thr Lys Thr Leu Glu Ile Asp Leu Thr Ser  
Pro Val Asn Gly Lys Tyr Ser Lys Val Ala Val Arg Gly Ser His Asp Asn Tyr Gly Thr Asn Leu Leu  
Thr Arg Asn Gly Ser Glu Trp Thr Gly Ile Phe Glu Phe Ser Lys Thr Thr Lys Phe Lys Leu Glu Ala  
Leu Pro Pro Ala Ala Asp Gln Cys Ile Phe Leu Gly Gly Asn Arg Gly Glu Ala Leu Thr Ala Ser Gly  
Gly Phe Ile Ser Leu Pro Ala Gly Arg Tyr Thr Ile Lys Phe Asn Glu Glu Ser Lys Val Leu Thr Ala  
Gly Asp Val Asp Cys Thr Gly

SEQ ID NO: 151

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tggtgaccacagtcggcctataccggcagcagtagctccatgggctacgaccgctgtattacttcgaccagcattcatcgttcggcaccgaag  
agcagctacggctcgttcacagtagctacacagcagaaaggaaactggcatcatagccgatgtagtgtcaatcaccgaaagaatgtctcaactg  
ggtggatttcccgccgagacctacaacgggtgaacctatcagatggtaagcaccgacatcgttcgaacgatgacggcggaaaaacagcca  
cttgggcaaatcaaacggctacagctctctctcaatgccgacgaaggcgaaggctgggacggcatgcgcgacctggaccacaagtgcga  
gaacgtgcagaaatcgggtcttgctacaccaaatactgtggtgacgacttaggctataccggattccgctacgatattggtaaaggatttgacgg  
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acagcaccaaaaagagcgtatgtccgcagtcggcagccttcgacttcgtttccgatacacctgccgcgatgccgtcaacaacaagaactgg  
gcgaacctgaagaacacttcgggtatcagcgtatccgattacaggcgtattcgggtacgtttgtgaaatcacgatacggaaatcaccgttcagct  
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accgactgcaaggagagatcaagaatctcatcgaggcagctgcctgtcgtgattcacaaccagagcacctatgccgaatggatgagcgg  
tgcagcctacatcgacgtaccgtaacaggtacgaacggcaccttacgtgttctgtcggctcttatcagtataatgtagccgccaactacattca  
gattctctcaggcaaaaactataatactacgtactcaacacgctcgaggctccttgatcgggaaaggttccggctctacaccgaaggtgaa  
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gccgtaaccagcgggaacggaactgaccatcacttcggacgccgtctgaaggttggtctgctttccggcgcatcgtcaggaaacatacagagc  
cgtacattcaccttcaggctgcaaacacctccgagtattacacagccaccatgcacgtatgcaaccagtcggagctctcaatccgctgtttgc  
ctatgttgggcaggaccggacaacgagcagattaacggcaactggccgggcaccaagctaccgctaccattaccgaaacaaccttacct

Figure 16 (cont.)

gggtacacgcagtcgtccagattccgaagaacgtggactatgtcgtgaactttgtttcaccacaaccggcggttacgcagacagtggatgtt  
accggcatgaaggccgatgtctgttacattattaacagtaccaagagcggcaacaagtacacggtaaccgacgttacctcacagtattctcgtt  
agaggccatctttgatgaagaaaactccggctccttccctgtctatgacctgcaggacgccgcgtcagcgaaattagaacaggacaattatat  
cttcagaacggaaagaagatactcatcagataaacagaggttccgaaccattctctattatgaaaatcagacacttagtaatctcagcactgctg  
ggtttggggggcctgtacaccatcagctgctcctcgtcggg

SEQ ID NO: 152

Met Lys Thr Ile Leu Ser Thr Ile Met Val Met Ala Ala Ala Ala Thr Thr Val Glu Ala Gln Gly  
Trp Pro Glu Asn Tyr Gly Gly Val Met Leu Gln Gly Phe Tyr Trp Asp Ser Tyr Ser Ala Thr Lys Trp  
Thr Lys Leu Glu Ala Gln Ala Asp Glu Ile Cys Asn Tyr Phe Ser Leu Val Trp Val Pro Gln Ser Ala  
Tyr Thr Gly Ser Ser Thr Ser Met Gly Tyr Asp Pro Leu Tyr Tyr Phe Asp Gln His Ser Ser Phe Gly  
Thr Glu Glu Gln Leu Arg Ser Phe Ile Ser Thr Tyr Lys Gln Lys Gly Thr Gly Ile Ile Ala Asp Val Val  
Val Asn His Arg Lys Asn Val Ser Asn Trp Val Asp Phe Pro Ala Glu Thr Tyr Asn Gly Val Thr Tyr  
Gln Met Val Ser Thr Asp Ile Val Ser Asn Asp Asp Gly Gly Lys Thr Ala Thr Trp Ala Asn Gln Asn  
Gly Tyr Ser Leu Ser Ser Asn Ala Asp Glu Gly Glu Gly Trp Asp Gly Met Arg Asp Leu Asp His  
Lys Ser Gln Asn Val Gln Lys Ser Val Leu Ala Tyr Thr Lys Tyr Leu Val Asp Asp Leu Gly Tyr Thr  
Gly Phe Arg Tyr Asp Met Val Lys Gly Phe Asp Gly Ser His Val Ala Asp Tyr Asn Thr Asn Ala  
Gly Val Gln Phe Ser Val Gly Glu Tyr Trp Asp Gly Thr Ala Ser Lys Val Tyr Ser Trp Ile Asn Ser  
Thr Lys Lys Ser Asp Val Pro Gln Ser Ala Ala Phe Asp Phe Ala Phe Arg Tyr Thr Cys Arg Asp Ala  
Val Asn Asn Lys Asn Trp Ala Asn Leu Lys Asn Thr Ser Gly Ile Ser Asp Ala Asp Tyr Arg Arg Tyr  
Ser Val Thr Phe Val Glu Asn His Asp Thr Glu Tyr Arg Ser Ala Thr Ala Ser Gln Asp Pro Ile Lys  
Gly Asp Thr Val Ala Leu Asn Ala Trp Met Leu Ala Met Pro Gly Thr Pro Cys Val Phe Leu Lys His  
Trp Thr Asp Cys Lys Glu Glu Ile Lys Asn Leu Ile Glu Ala Arg Arg Leu Val Gly Ile His Asn Gln  
Ser Thr Tyr Ala Glu Trp Met Ser Gly Ala Ala Tyr Ile Gly Arg Thr Val Thr Gly Thr Asn Gly Thr  
Leu Arg Val Leu Cys Gly Ser Tyr Gln Tyr Asn Val Ala Ala Asn Tyr Ile Gln Ile Leu Ser Gly Lys  
Asn Tyr Lys Tyr Tyr Val Leu Asn Thr Leu Glu Ala Pro Trp Ile Gly Lys Gly Ser Gly Ser Tyr Thr  
Glu Gly Glu Thr Val Thr Val Pro Leu Ile Ala Ile Ser Ala Asp Ala Asn Ala Lys Leu Val Tyr Thr  
Thr Asp Gly Thr Asp Pro Thr Ala Thr Ser Thr Ala Val Thr Ser Gly Thr Glu Leu Thr Ile Thr Ser  
Asp Ala Val Leu Lys Val Gly Leu Leu Ser Gly Gly Ile Val Arg Asn Ile Gln Ser Arg Thr Phe Thr  
Phe Gln Ala Ala Asn Thr Ser Glu Tyr Tyr Thr Ala Thr Met His Val Cys Asn Gln Ser Gly Ala Leu  
Asn Pro Leu Phe Ala Tyr Val Trp Ala Gly Pro Asp Asn Glu Gln Ile Asn Gly Asn Trp Pro Gly Thr  
Lys Leu Thr Ala Thr Ile Thr Glu Asn Asn Leu Thr Trp Tyr Thr Gln Ser Phe Gln Ile Pro Lys Asn  
Val Asp Tyr Val Val Asn Phe Val Phe Thr Thr Thr Gly Gly Gly Thr Gln Thr Val Asp Val Thr Gly  
Met Lys Ala Asp Val Trp Tyr Ile Ile Asn Ser Thr Lys Ser Gly Asn Lys Tyr Thr Val Thr Asp Val  
Thr Ser Gln Tyr Ser Ser Leu Glu Ala Ile Phe Asp Glu Glu Asn Ser Gly Ser Phe Pro Val Tyr Asp  
Leu Gln Gly Arg Arg Val Ser Glu Ile Arg Asn Arg Thr Ile Ile Ser Ser Glu Arg Lys Glu Asp Thr  
His Gln Ile Asn Arg Gly Ser Glu Pro Phe Ser Tyr Tyr Glu Asn Gln Thr Leu Ser Asn Leu Ser Thr  
Ala Gly Phe Gly Gly Leu Val His His Gln Leu Leu Leu Val Gly

SEQ ID NO: 69

atgttgaaaaggattacggtagtctgtttattgttttttgccttttctaatatatatgagggaaataaggcagaagcagcaacagtgaacaatgga  
acattaatgcagtattttgagtgttacgctccgaatgatgggaatcattggaatcggttgcgttccgatgctgaaagttagctcataaaggaatcac  
atctgtatggataccacctgcataaaagggaacttcgcaaaatgatgtagggtaggggcctatgatttatatgatttaggggagttcaatcaaaaa  
ggaacggtgcggacgaaatatgggacaaaagcacagttgaaatctgcaattgacgctttacataagcaaaacatcgacgtatacgggtgatgtag  
ttatgaatcataaagggtggggctgattatactgaaaccgtaacagctgttgaggtagaccgtaacaatcgaaatattgaagatcaggtgattatca  
aattagtgcattggacgggggttaattttccaggggcgcggagatgctattctaatttcaaatggaaatgggtatcattttgacgggaacggattgggatg  
aagggaaggaaattaaatcgaattataaatttaggggtgtagataaagcgtgggattgggaagtgtctagcgaaatggaaattatgattttgat  
gtatgcagatcttgattttgatcatcctgatgttgcaatgagatgaaaaattgggaacatggtatgcgaatgaattaaattagatggcttcgtt  
ggacgctgtaaacatattgatcatgaattttacgcgattgggtaaatcatgccagacagcaaacggggaagaaatgttacagtagctgaata  
ttggcaaatgatgttcaggctttaacaattatttagcgaaagtcaattataatcaatctgtgtttgatgcaccgcttcattacaatttcattatgcttc

Figure 16 (cont.)

aacaggaaatgggaattatgatatgagaaatattttaaatggaaacagtaatgaaaaatcacctgcactgcagttactctcgttgagaatcatgat  
tctcagcctgggcagtcattggaatctgtagtaagtcctggtttaagccgctggcatatgcatttattttaactcgtgcagagggtatccttcagtt  
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attactcggacggaccaggagatcaaagtggatggatgttgaaagaataacgctggggaagtatggtacgatattacgggtaatacaaac  
aaatactgtaacaattaataaggacggatgggggcagttctatgtaagtggcggctcagttccatatatgttcagggtaa

SEQ ID NO: 70

Met Leu Lys Arg Ile Thr Val Val Cys Leu Leu Phe Ile Leu Leu Phe Pro Asn Ile Tyr Glu Gly Asn  
Lys Ala Glu Ala Ala Thr Val Asn Asn Gly Thr Leu Met Gln Tyr Phe Glu Trp Tyr Ala Pro Asn Asp  
Gly Asn His Trp Asn Arg Leu Arg Ser Asp Ala Glu Ser Leu Ala His Lys Gly Ile Thr Ser Val Trp  
Ile Pro Pro Ala Tyr Lys Gly Thr Ser Gln Asn Asp Val Gly Tyr Gly Ala Tyr Asp Leu Tyr Asp Leu  
Gly Glu Phe Asn Gln Lys Gly Thr Val Arg Thr Lys Tyr Gly Thr Lys Ala Gln Leu Lys Ser Ala Ile  
Asp Ala Leu His Lys Gln Asn Ile Asp Val Tyr Gly Asp Val Val Met Asn His Lys Gly Gly Ala Asp  
Tyr Thr Glu Thr Val Thr Ala Val Glu Val Asp Arg Asn Asn Arg Asn Ile Glu Val Ser Gly Asp Tyr  
Gln Ile Ser Ala Trp Thr Gly Phe Asn Phe Pro Gly Arg Gly Asp Ala Tyr Ser Asn Phe Lys Trp Lys  
Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Glu Gly Arg Lys Leu Asn Arg Ile Tyr Lys Phe Arg Gly  
Val Asp Lys Ala Trp Asp Trp Glu Val Ser Ser Glu Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp  
Leu Asp Phe Asp His Pro Asp Val Ala Asn Glu Met Lys Asn Trp Gly Thr Trp Tyr Ala Asn Glu  
Leu Asn Leu Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Asp His Glu Tyr Leu Arg Asp Trp Val  
Asn His Ala Arg Gln Gln Thr Gly Lys Glu Met Phe Thr Val Ala Glu Tyr Trp Gln Asn Asp Val Gln  
Ala Leu Asn Asn Tyr Leu Ala Lys Val Asn Tyr Asn Gln Ser Val Phe Asp Ala Pro Leu His Tyr Asn  
Phe His Tyr Ala Ser Thr Gly Asn Gly Asn Tyr Asp Met Arg Asn Ile Leu Asn Gly Thr Val Met Lys  
Asn His Pro Ala Leu Ala Val Thr Leu Val Glu Asn His Asp Ser Gln Pro Gly Gln Ser Leu Glu Ser  
Val Val Ser Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Ala Glu Gly Tyr Pro Ser  
Val Phe Tyr Gly Asp Tyr Tyr Gly Thr Ser Gly Asn Ser Ser Tyr Glu Ile Pro Ala Leu Lys Asp Lys  
Ile Asp Pro Ile Leu Thr Ala Arg Lys Asn Phe Ala Tyr Gly Thr Gln Arg Asp Tyr Leu Asp His Pro  
Asp Val Ile Gly Trp Thr Arg Glu Gly Asp Gly Val His Ala Asn Ser Gly Leu Ala Thr Leu Leu Ser  
Asp Gly Pro Gly Gly Ser Lys Trp Met Asp Val Gly Lys Asn Asn Ala Gly Glu Val Trp Tyr Asp Ile  
Thr Gly Asn Gln Thr Asn Thr Val Thr Ile Asn Lys Asp Gly Trp Gly Gln Phe Tyr Val Ser Gly Gly  
Ser Val Ser Ile Tyr Val Gln Arg

SEQ ID NO: 153

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atgagcctatcaacaacatacaagcagaaacattacataataaagggtcaaaaggcgcaaacaggaaataaagacggaatttttatgaact  
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cagtatggaagtttacaagatttccgtgaactaacaacagaagcgcataaacgcaacgtaaggtagtaatagatcttggttataatcatacaagc  
agtgagcatccttggttgtcgatgcataaaaaataaaaacagtaagtagcagattactatatttgggctgataaaaatacagactaaatgaaa  
aggcccatggggcaacaagtatggcacaagcgtcgaacggagagatatttctacgcaacgcttctgggaagggatgccggacttaaatatga  
caacctaaagtaagagaagaatgattaaaatcgggaaattttggctcaacaaggagctgatggcttctgctatagtcagccatgcacatctt  
taaggggcaaacacctgaaggagcaaaagaaaaatattgaatggtggaatgaattccgcgacgcgatgagagaaacgaatccaaatacgtatct  
agttggtgaaatatgggatcaaccagaagtagttgctccgtattatcaatcgttagattctacatttaacttcgacttagcatataaaatcgttaattcc  
gttaaaaatggtactgatcaaggggtagccgcggcagctgttgcaacggatgagttatataaaacataataatccaaataaaattgatggaacgttt  
ttaacgaatcatgacaaaatcgtgtaatgagtgagttaaatggtgatgtaacaaagcaaaatcagcagcctctattctgttgacactccctggta  
atccgttcatttattatggcgaagaaatcggcatgacaggccaaaacagatgagttgattcgtgagccttccgttggtatgaagatgataaag  
aagggtcaaacgagctgggagactccagtatataacattgatcataatggtgtttcagttgaagcacaagataaacaaggtctctcttaagcc  
attatcgtaaaatgattcgtgttcgtagcaacacgatgaactgtcaaaaggtaatttagaacctatttctgtcaataattcacaggttggtgcataat  
cgtacgtataaaaaataatcaattcaagtgtaccataatatttcagacaagccggttacattaactgtttcaacaaggaaggaactgatttttctagt  
gaattaggagcaaaaaggaaaaatcaacattagtaattccagcgaatacgcagtgctagtaagtaa

Figure 16 (cont.)

SEQ ID NO: 154

Met Pro Ser Ile Asn Ala Ser Asp Cys Lys Lys Lys Gly Asp Arg Ser Met Lys Arg Lys Lys Trp Thr  
Ala Leu Ala Leu Ser Leu Pro Leu Val Met Ser Leu Ser Thr Asn Ile Gln Ala Glu Thr Leu His Asn  
Asn Lys Gly Gln Lys Ala Gln Thr Gly Asn Lys Asp Gly Ile Phe Tyr Glu Leu Tyr Val Asn Ser Phe  
Tyr Asp Thr Asp Ser Asn Gly His Gly Asp Leu Lys Gly Val Thr Lys Lys Leu Asp Tyr Leu Asn  
Asp Gly Asn Pro Arg Thr Asn Asn Asp Leu Gln Ile Asn Gly Ile Trp Met Met Pro Ile Asn Thr Ser  
Pro Ser Tyr His Lys Tyr Asp Val Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Ser Leu Gln Asp  
Phe Arg Glu Leu Thr Thr Glu Ala His Lys Arg Asn Val Lys Val Val Ile Asp Leu Val Ile Asn His  
Thr Ser Ser Glu His Pro Trp Phe Val Asp Ala Leu Lys Asn Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr  
Ile Trp Ala Asp Lys Asn Thr Asp Leu Asn Glu Lys Gly Pro Trp Gly Gln Gln Val Trp His Lys Ala  
Ser Asn Gly Glu Tyr Phe Tyr Ala Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Lys  
Val Arg Glu Glu Met Ile Lys Ile Gly Lys Phe Trp Leu Lys Gln Gly Ala Asp Gly Phe Arg Leu Asp  
Ala Ala Met His Ile Phe Lys Gly Gln Thr Pro Glu Gly Ala Lys Lys Asn Ile Glu Trp Trp Asn Glu  
Phe Arg Asp Ala Met Arg Glu Thr Asn Pro Asn Thr Tyr Leu Val Gly Glu Ile Trp Asp Gln Pro Glu  
Val Val Ala Pro Tyr Tyr Gln Ser Leu Asp Ser Thr Phe Asn Phe Asp Leu Ala Tyr Lys Ile Val Asn  
Ser Val Lys Asn Gly Thr Asp Gln Gly Val Ala Ala Ala Val Ala Thr Asp Glu Leu Tyr Lys Thr  
Tyr Asn Pro Asn Lys Ile Asp Gly Thr Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu  
Asn Gly Asp Val Asn Lys Ala Lys Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Phe Ile  
Tyr Tyr Gly Glu Glu Ile Gly Met Thr Gly Gln Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp  
Tyr Glu Asp Asp Lys Glu Gly Gln Thr Ser Trp Glu Thr Pro Val Tyr Asn Ile Asp His Asn Gly Val  
Ser Val Glu Ala Gln Asp Lys Gln Lys Ala Ser Leu Leu Ser His Tyr Arg Lys Met Ile Arg Val Arg  
Gln Gln His Asp Glu Leu Val Lys Gly Asn Leu Glu Pro Ile Ser Val Asn Asn Ser Gln Val Val Ala  
Tyr Asn Arg Thr Tyr Lys Asn Lys Ser Ile Gln Val Tyr His Asn Ile Ser Asp Lys Pro Val Thr Leu  
Thr Val Ser Asn Lys Gly Lys Leu Ile Phe Ser Ser Glu Leu Gly Ala Lys Lys Glu Lys Ser Thr Leu  
Val Ile Pro Ala Asn Thr Thr Val Leu Val Lys

SEQ ID NO: 155

gtgtcaagaatgtttgcaaacgattcaaacctctttactgccgttatcgtggtttttattgctgtttcatttggttctggcaggaccaacggctg  
cgaatgctgaacgggctaacaatcaaatgagcttacagcaccgtcgatcaaaagcgggaaccattctcatgcttggaaatggctgccaatacgt  
taaacacacaatatgaaggatattcatgatgcaggatatacagcgattcagacgtctccgattaaccaagtcaaggaagggaaccaaggaaataa  
aaacatgtcgaactgggtactggctctatcagccgacatcgatccaaattggcaaccgttacttaggtactgaacaagaattaaagaatgtgtgc  
agccgctgaagaatatggcataaagggtattgttgacgcggatcatcaatcataccaccagtactatgccgcatgttccaatgagattaagagtatt  
ccaaactggacacatggaaacacacaaataaaaaactgggtctgatcgatgggatgtcacgcagaatgcattgctcgggctgtatgactggaata  
cacaaaatacacaaagtacagtcctatttgaacgggtcttagaaagagcattgaatgacggggcagacgggtttcgatttgatgccgccaacata  
tagagcttccggatgatggcagttacggcagtcattttggccgaatatcacaaatcatctgcagagttccaatacggagaaatcctgcaggat  
agtgttcaagagatgcttcattatgcgaattatatgaatgtgacagcgtctaactatgggcattccataaggctccgtttaaagaatcgaatctggg  
cgtgtcgaatatctcccactatgcatcagatgtgtctgcggacaagctagtacatgggtagaatcgcatgatacgtatgccaatgatgatgaag  
agtcgacatggatgagcgtatgatatacgttttaggctgggctgtagcttctcgttcaggcagtagcgcctcttttcttccagacctgaggg  
aggcggaaatggtgtgagattcccgggaaaagccaaataggcgatcgcgaggatgctttattgaagatcaggctatcactgcgggtcaatag  
attcacaaatgtgatggctggacagcctgaggaactctgaacccaaatggaaacaaccagatatattgaatcagcgcggtcacatggcggtg  
tgctggcaaatgcaggttcacctctgtttctatcaatacgcacaaatggcctgatggcaggtatgataataaagctggggcaggttcattca  
agtaaatgacggtaaaactgacaggcacgatcaatgccaggtctgtggctgtgctttatcctgatgatattgcaaaagcgctcatgtttccttgag  
aattacaaaacaggtgtaacacattcttcaatgatcaactgacgattacactgcgtgcagatgcgaatacaacaaaagcggttatcaaatcaata  
atggaccagagacggcggttaaggatggagatcaattcacaatcgaaaaggagatccatttgcaaaacatacaccatcatgttaaaagggaac  
gaacagtgtgtgtaacgaggaccgaggaatacagttttgtaaaagagatccagcttcggccaaaaccatcggtatcaaaatccgaatcatt  
ggagccaggtaaatgcttatatctataaacatgatggggggccgggca

SEQ ID NO: 156

Figure 16 (cont.)

Val Ser Arg Met Phe Ala Lys Arg Phe Lys Thr Ser Leu Leu Pro Leu Phe Ala Gly Phe Leu Leu Leu  
Phe His Leu Val Leu Ala Gly Pro Thr Ala Ala Asn Ala Glu Thr Ala Asn Lys Ser Asn Glu Leu Thr  
Ala Pro Ser Ile Lys Ser Gly Thr Ile Leu His Ala Trp Asn Trp Ser Phe Asn Thr Leu Lys His Asn  
Met Lys Asp Ile His Asp Ala Gly Tyr Thr Ala Ile Gln Thr Ser Pro Ile Asn Gln Val Lys Glu Gly  
Asn Gln Gly Asn Lys Asn Met Ser Asn Trp Tyr Trp Leu Tyr Gln Pro Thr Ser Tyr Gln Ile Gly Asn  
Arg Tyr Leu Gly Thr Glu Gln Glu Phe Lys Glu Met Cys Ala Ala Ala Glu Glu Tyr Gly Ile Lys Val  
Ile Val Asp Ala Val Ile Asn His Thr Thr Ser Asp Tyr Ala Ala Ile Ser Asn Glu Ile Lys Ser Ile Pro  
Asn Trp Thr His Gly Asn Thr Gln Ile Lys Asn Trp Ser Asp Arg Trp Asp Val Thr Gln Asn Ala Leu  
Leu Gly Leu Tyr Asp Trp Asn Thr Gln Asn Thr Gln Val Gln Ser Tyr Leu Lys Arg Phe Leu Glu  
Arg Ala Leu Asn Asp Gly Ala Asp Gly Phe Arg Phe Asp Ala Ala Lys His Ile Glu Leu Pro Asp Asp  
Gly Ser Tyr Gly Ser Gln Phe Trp Pro Asn Ile Thr Asn Thr Ser Ala Glu Phe Gln Tyr Gly Glu Ile  
Leu Gln Asp Ser Ala Ser Arg Asp Ala Ser Tyr Ala Asn Tyr Met Asn Val Thr Ala Ser Asn Tyr Gly  
His Ser Ile Arg Ser Ala Leu Lys Asn Arg Asn Leu Gly Val Ser Asn Ile Ser His Tyr Ala Ser Asp  
Val Ser Ala Asp Lys Leu Val Thr Trp Val Glu Ser His Asp Thr Tyr Ala Asn Asp Asp Glu Glu Ser  
Thr Trp Met Ser Asp Asp Asp Ile Arg Leu Gly Trp Ala Val Ile Ala Ser Arg Ser Gly Ser Thr Pro  
Leu Phe Phe Ser Arg Pro Glu Gly Gly Gly Asn Gly Val Arg Phe Pro Gly Lys Ser Gln Ile Gly Asp  
Arg Gly Ser Ala Leu Phe Glu Asp Gln Ala Ile Thr Ala Val Asn Arg Phe His Asn Val Met Ala Gly  
Gln Pro Glu Glu Leu Ser Asn Pro Asn Gly Asn Asn Gln Ile Phe Met Asn Gln Arg Gly Ser His Gly  
Val Val Leu Ala Asn Ala Gly Ser Ser Ser Val Ser Ile Asn Thr Pro Thr Lys Leu Pro Asp Gly Arg  
Tyr Asp Asn Lys Ala Gly Ala Gly Ser Phe Gln Val Asn Asp Gly Lys Leu Thr Gly Thr Ile Asn Ala  
Arg Ser Val Ala Val Leu Tyr Pro Asp Asp Ile Ala Lys Ala Pro His Val Phe Leu Glu Asn Tyr Lys  
Thr Gly Val Thr His Ser Phe Asn Asp Gln Leu Thr Ile Thr Leu Arg Ala Asp Ala Asn Thr Thr Lys  
Ala Val Tyr Gln Ile Asn Asn Gly Pro Glu Thr Ala Phe Lys Asp Gly Asp Gln Phe Thr Ile Gly Lys  
Gly Asp Pro Phe Gly Lys Thr Tyr Thr Ile Met Leu Lys Gly Thr Asn Ser Asp Gly Val Thr Arg Thr  
Glu Glu Tyr Ser Phe Val Lys Arg Asp Pro Ala Ser Ala Lys Thr Ile Gly Tyr Gln Asn Pro Asn His  
Trp Ser Gln Val Asn Ala Tyr Ile Tyr Lys His Asp Gly Gly Arg Ala

SEQ ID NO: 157

atgcaaacgattgcaaaaaaggggatgaacgatgaaagggaataatggacagcattagcttaacactgccgctggctgctagcttatca  
acaggcggtcacgccgaacccgtacataaaggtaaacgtccaacagcagataaaacgggtgtctttatgaggtgtatgtaaactcttttacgat  
gcaataaagatggacatggtgatttaaaagggtcttacacaaaagctggattttgaatgacggcaattctcatacaaaatgatcttcaagtaa  
acggaatttgatgatgccggtaaacctctcctagctatcataatatgatgtaacggactattataacattgatccgcagtacggaaatctgca  
agattttcgcaagctgatgaaagaagcagataaacgagacgtaaaagggtattatggacctcgttgtaatacatacagcagtgaaatccttggtt  
tcaagctgcattaaaagataaaaacagcaagtacagagattactatatttggccgataaaaatactgatttaaatgaaaaaggatcttgggggca  
gcaagtatggcataaaagctccaaacggagagatattttatggtacgttttgggaaggaaatgcctgacttaaatcagataatcccgaagtaagaaa  
agaaatgattaacgtcgggaaattttggctaaagcaaggcgttgacgggtccgcttagatgctgcgttcataatttttaagggtcaaacacctgaa  
ggcgctaagaaaaatcgtgtggtggaatgagtttagagatgcaatgaaaaagaaaaccctaactatatacaggggtgaagtatgggatc  
aaccggaagtagtagctccttactatcaatcgttgattctttttaactttgatttagcaggaaagattgtaaactctgtaaaatcaggaaatgatca  
aggaatcgcgactgcagcagccgcaactgatgagctgttcaaatcatacaatccaaataaaatgacggcattttctaaccaacctgacccaaa  
atcgcgtcatgagtgcagtaagcggcgatgtgaataaagcaaaagtcagctgccttatcttacttacgcttcctggcaacccgatatatttacgg  
tgaagaaattggaatgaccgggtgaaaagcctgatgagttatccgtgaaccgttccgctggtacgaaggcaatggacttggacaaaccagctg  
ggaaacatccgtatacaacaaaggcggcaatggtgtgtcagtagagacacaaacaaaaggattctttgttaaatcattaccgtgaaatga  
ttcgctgctcagcagcatgaagagttagtaaaaggaaccttcaatctatttcagtagacagtaaagaagtcgttgccatagccgcacgtata  
aaggcaaatcgattagcgtgtatcataatatttcaaatcaaccggtaaaagtaatctgtaacagcgaaaggtaaatgatttttgctagtgaaaaagg  
gcaaaaaagtcaaaaatcagcttgggttcagctaatacaacggttttaataaaaaa

SEQ ID NO: 158

Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr  
Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ala Pro Thr  
Ala Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His

Figure 16 (cont.)

Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn  
Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val  
Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala  
Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe  
Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp  
Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly  
Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val  
Gly Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly  
Gln Thr Pro Glu Gly Ala Lys Lys Asn Ile Val Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu  
Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser  
Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Asn Ser Val Lys Ser Gly Asn Asp Gln  
Gly Ile Ala Thr Ala Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly  
Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala Lys  
Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr  
Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr  
Ser Trp Glu Thr Ser Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Thr Gln Thr Lys Gln Lys  
Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly  
Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Lys Ser Ile  
Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Thr Ala Lys Gly Lys Leu Ile Phe Ala  
Ser Glu Lys Gly Ala Lys Lys Val Lys Asn Gln Leu Val Val Pro Ala Asn Thr Thr Val Leu Ile Lys

SEQ ID NO: 159

ttgcaaaaaaagggatgaaacgatgaaagggaataatggacagctttagctctaactgccgtggctgctagcttatcaacaggcggttc  
acgccgaaaccgtacataaaggtaaatctccaacagcagataaaaacgggtgattttatgaggtgatgtaactcttttacgatgcaataaaga  
tggacatgggtgatttaaaaggcttacacaaaagggtgattttaaatgatggcaattctcatacaagaatgatcttcaagtaaacgggatttggat  
gatgccggtcaacccttctccagctatcataaatatgatgtaacggactattataatattgatccgcagatggaaatctgcaagatttgcgaaac  
tgatgaaagaagcagataaacgagatgtaaaagtcattatggacctcgttgtaacatacagcagtgaaacacccttgggttcaagctgcattaa  
aagataaaaacagcaagtacagagattactatctgggctgataaaaataccgactgaatgaaaaggatcttggggacagcaagtatggca  
taaagctccaaacggagagtattttacggaacggtttgggaagggaatgccggacttaattacgataatcctgaagtaagaaaagaaatgattaa  
cgtaggaaagtttggctaaagcaaggagttgatgggtccgcttagatgctgcgttcataatttttaaggccaaacacctgaaggcgtaagaa  
aatctcctgtggtggaatgaatttagagatgcaatgaaaaaggaaaacctaactatataacgggtgaagtatgggatcaaccggaagta  
gtagctccttactatcaatcgcttgattctttatctaatttgatttagcaggaaagattgtaactctgtaaaatcaggaaatgatcaaggaatcgga  
ctgcagcagcggaacggatgaactgttcaaatcatacaatccaaataaaaattgacgggtattttctaaccaaccatgacaaaatcgctcatga  
gtgagctaaacggcgatgtgaataaagcaagtcagctgccttacttactacgcttcctggcaacccgtatatttattacgggtgaagaaatcgga  
catgaccggtgaaaagcctgatgagtaatccgtgaaccgttccctggtagcaaggaaacggacttggacaaaccagctgggaaacacctgt  
atatacaaaaggcggcaacggcgtgctgtagaagcacaacaaaacaaaaggactcttgttaaatcattaccgtgaaatgattcgcgtgcgtc  
agcagcacgaagagttagtaaaaggaacgcttcaatctatttcagtagacagtaaaagtcgttgctatagccgtacgtataaaggcaaatcg  
attagcgtgtatcataatatttcaaatcaaccggtaaaagtatctgtagcagcaaaaggtaaatgattttgctagtgaaggggtgctaagaaagt  
caaaaatcagcttgtgattccggcgaatacaacggttttaataaaaataa

SEQ ID NO: 160

Met Gln Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr Leu Pro  
Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ser Pro Thr Ala Asp  
Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His Gly Asp  
Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn Asp Leu  
Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val Thr Asp  
Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala Asp  
Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe Gln  
Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp Leu  
Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly Thr

Figure 16 (cont.)

Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val Gly  
Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly Gln  
Thr Pro Glu Gly Ala Lys Lys Asn Leu Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu  
Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser  
Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Asn Ser Val Lys Ser Gly Asn Asp Gln  
Gly Ile Ala Thr Ala Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly  
Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Asn Gly Asp Val Asn Lys Ala Lys  
Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr  
Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Pro Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr  
Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln Lys  
Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly  
Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Lys Ser Ile  
Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile Phe Ala  
Ser Glu Lys Gly Ala Lys Lys Val Lys Asn Gln Leu Val Ile Pro Ala Asn Thr Thr Val Leu Ile Lys

SEQ ID NO: 161

gtggatccaaagaattgtatgcaatttatgcaaacgattgcaaaaaaggggatgaacgatgaaagggaaaaatggacagcttttagctctaa  
cactgccgtggtgctagcttatcaacaggtgttcacgccgaaccgtacataaaggtaaagctccaacagcagataaaaaagggtgtctttat  
gaggtatatgtaaaactcttttacgatgcaataaagatggacatggtgatttaaaaggccttacacaaaaggtagactatataatgacggaaattc  
tcatacaaaagaatgatcttcaagtaaacgggatttgatgatgccgtcaacccctctcctagctatcataaatatgatgtaacggactattataatat  
tgatccgcagtagtgaatctgcaagatttgcgaacttatgaaagaagcagataaacgagacgtaaaagtcattatggaccttgtgtgaatcat  
acgagcagtgaaacacccctgtttcaagctgcgttgaaagataaaaacagcaagtacagagattactatatttgggctgataaaaatactgacttg  
aatgaaaaaggatcttggggacaacaagtatggcataaagctccaaacggagagtagtttttacggaacgttctgggaaggaatgcctgacttaa  
attacgataaccctgaagtaagaaagaaatgattaacgtcggaaagtgttgctaaaacaaggcgttgacggctccgcttagatgctgcccttc  
atatttttaaggtaaacgcctgaaggcgtaagaaaaacattctatggtggaatgagtttagagatgcatgaaaaagaaaaccgaacgta  
tatctaacgggtgaagtgtggaccagccagaagtagtagcccttactatcaatcacttgattctctatttaatttgatttagcaggaaaaattgtc  
agctctgtaaaagcaggaaatgatcaaggaatcgccactgcagcagcggaactgatgagctgttcaaatcatacaatccaaataaaattgacg  
gcattttcttaaccaaccatgaccaaatacgcgtcatgagtgagtgtaagcggcgatgtgaataaagcaaaatcagccgctctatcttacttacgct  
tcttggaatccgtatatttattacgggtgaagaattggcatgacaggtgaaaagcctgatgaattaatccgtgaaccgttccgctggtacgaagg  
caacggaattggacaaactagctgggaaacacctgtatatacaaaggcggtaacggcggtgtctgtagaagcacaacaaaacaaaaggatt  
cctgttaaatcattaccgtgaaatgattcgtgtgcgcagcagcagcaagagtagtaaaaaggaacgcttcaatccatttcagtagacagtaaaag  
aagtcgttcctatagccgcacgtacaaaggcaaatcgattagcgtgtatcataatatttcaaatcaacctgtaaaagtagtctgtagcagcgaaag  
gtaacttgattttgtagtgaaaaagggtgtaagaaagtcaaaaatcagcttgattccggcgaatgcgacgggtttaataaaaataa

SEQ ID NO: 162

Val Asp Pro Lys Asn Cys Ser Gln Phe Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly  
Lys Lys Trp Thr Ala Leu Ala Leu Thr Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu  
Thr Val His Lys Gly Lys Ala Pro Thr Ala Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser  
Phe Tyr Asp Ala Asn Lys Asp Gly His Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu  
Asn Asp Gly Asn Ser His Thr Lys Asn Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro  
Ser Pro Ser Tyr His Lys Tyr Asp Val Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Asn Leu Gln  
Asp Phe Arg Lys Leu Met Lys Glu Ala Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val  
Asn His Thr Ser Ser Glu His Pro Trp Phe Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp  
Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His  
Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp  
Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val Gly Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe  
Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly Gln Thr Pro Glu Gly Ala Lys Lys Asn Ile Leu Trp  
Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp  
Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly  
Lys Ile Val Ser Ser Val Lys Ala Gly Asn Asp Gln Gly Ile Ala Thr Ala Ala Ala Ala Thr Asp Glu

Figure 16 (cont.)

Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val  
Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala Lys Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly  
Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu  
Pro Phe Arg Trp Tyr Glu Gly Asn Gly Ile Gly Gln Thr Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly  
Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln Lys Asp Ser Leu Leu Asn His Tyr Arg Glu Met  
Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly Thr Leu Gln Ser Ile Ser Val Asp Ser Lys  
Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Lys Ser Ile Ser Val Tyr His Asn Ile Ser Asn Gln Pro  
Val Lys Val Ser Val Ala Ala Lys Gly Asn Leu Ile Phe Ala Ser Glu Lys Gly Ala Lys Lys Val Lys  
Asn Gln Leu Val Ile Pro Ala Asn Ala Thr Val Leu Ile Lys

SEQ ID NO: 163

atggtacgtcccgaaacgacgggctgattggaaccgactatcgaacgactcgcagcacttgaaagacattgggtgacgacgggtgtgattccg  
ccggcgctacaaaggcagtcacagaaacgatgtcgggtatggggcgctacgatttatacgcctcggcgaattcaacaaaaaggacgacccg  
gacgaagtacgggacgaaagcgcagctccagaccgcatctcgaacttgcgcggtaaaggatcggtgtgtacggcgacgtcgtcatgaat  
cacaagggcgggggcattataccgaatccgttcaggcgatcgagggtcaatccgtcgaaccggaaccaagaacgtccggtgagtatggcat  
ctcggcctggactgggttcaacttcgcggggcgcaacaatacatactcgccttcaaatggcgctggtaccattttgacggtagcattgggac  
agtcacgcagcttgagccgcactctataagttcaagagcacaggcaaggcggtgggacacggacgtgtcgaacgagaacggcaactatgattat  
cttatgtatgccgacgtcgatttcgagcatcccgaggtccgccaagagatgaagaactggggcgaatggtacgccgactcgtcgggctcgac  
ggtttccgggtggatgcggtcaaacatcagccactcgtactgaaggagtggtgacgagcgtgcgccagacgacgggaaagagatgttc  
acggtcgccgagtttggaagaacgatctcggtgccatcaacgactatctgtataagacgggctacacgactccgtcttcgatgtccgctcc  
attataactccaagcggcggttaacggcgggcggtattacgatatgcgcaacatcttgaaggcaccgtcaccgaacagcatccgtcgtcgtc  
cgtgacgattgtcgataaccacgactcacagccggcgagtcgctcgagtcgacggtcgccaactggttcaaacgctcgcctacgcgacga  
tcacgacgcgggtcagggttatccggccctcttctatggagactattatggcacgaaagggacgacgaaccgcgaatccgaacatgtcgg  
gcacgctccaaccgattttgaaggcacgaaaagacttcgcctacgggacgcagcatgactacctcgaatcagggacgtcgttgacac  
gtgaaggtgtgaccgaccgtgccaaatcggtctcgcgacgattctatcgacgggtccggcggtcgaagtggatgtacgtcggcaaacag  
aacgccggcgaggtatgaaagacatgacgaacaacaacgcccgtctcgtcacgatcaatgctgacgggtggggtcagttcttcgtaacgg  
aggctcgggtctcgatttatacgaacaataa

SEQ ID NO: 164

Met Val Arg Pro Glu Arg Arg Ala Ala Leu Glu Pro Thr Ile Glu Arg Leu Ala Ala Leu Glu Arg His  
Trp Val Thr Thr Val Trp Ile Pro Pro Ala Tyr Lys Gly Thr Ser Gln Asn Asp Val Gly Tyr Gly Ala  
Tyr Asp Leu Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Thr Arg Thr Lys Tyr Gly Thr Lys  
Ala Gln Leu Gln Thr Ala Ile Ser Asn Leu Arg Gly Lys Gly Ile Gly Val Tyr Gly Asp Val Val Met  
Asn His Lys Gly Gly Ala Asp Tyr Thr Glu Ser Val Gln Ala Ile Glu Val Asn Pro Ser Asn Arg Asn  
Gln Glu Thr Ser Gly Glu Tyr Gly Ile Ser Ala Trp Thr Gly Phe Asn Phe Ala Gly Arg Asn Asn Thr  
Tyr Ser Pro Phe Lys Trp Arg Trp Tyr His Phe Asp Gly Thr Asp Trp Asp Gln Ser Arg Ser Leu Ser  
Arg Ile Tyr Lys Phe Lys Ser Thr Gly Lys Ala Trp Asp Thr Asp Val Ser Asn Glu Asn Gly Asn Tyr  
Asp Tyr Leu Met Tyr Ala Asp Val Asp Phe Glu His Pro Glu Val Arg Gln Glu Met Lys Asn Trp  
Gly Lys Trp Tyr Ala Asp Ser Leu Gly Leu Asp Gly Phe Arg Leu Asp Ala Val Lys His Ile Ser His  
Ser Tyr Leu Lys Glu Trp Val Thr Ser Val Arg Gln Thr Thr Gly Lys Glu Met Phe Thr Val Ala Glu  
Tyr Trp Lys Asn Asp Leu Gly Ala Ile Asn Asp Tyr Leu Tyr Lys Thr Gly Tyr Thr His Ser Val Phe  
Asp Val Pro Leu His Tyr Asn Phe Gln Ala Ala Gly Asn Gly Gly Gly Tyr Tyr Asp Met Arg Asn Ile  
Leu Lys Gly Thr Val Thr Glu Gln His Pro Ser Leu Ser Val Thr Ile Val Asp Asn His Asp Ser Gln  
Pro Gly Gln Ser Leu Glu Ser Thr Val Ala Asn Trp Phe Lys Pro Leu Ala Tyr Ala Thr Ile Met Thr  
Arg Gly Gln Gly Tyr Pro Ala Leu Phe Tyr Gly Asp Tyr Tyr Gly Thr Lys Gly Thr Thr Asn Arg Glu  
Ile Pro Asn Met Ser Gly Thr Leu Gln Pro Ile Leu Lys Ala Arg Lys Asp Phe Ala Tyr Gly Thr Gln  
His Asp Tyr Leu Asp His Gln Asp Val Ile Gly Trp Thr Arg Glu Gly Val Thr Asp Arg Ala Lys Ser  
Gly Leu Ala Thr Ile Leu Ser Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln Asn Ala  
Gly Glu Val Trp Lys Asp Met Thr Asn Asn Asn Ala Arg Leu Val Thr Ile Asn Ala Asp Gly Trp Gly  
Gln Phe Phe Val Asn Gly Gly Ser Val Ser Ile Tyr Thr Gln Gln

SEQ ID NO: 165

Variable	Mean	SD	Min	Max	Median	Q1	Q3	Mode	Skewness	Kurtosis	Shapiro-Wilk	Normality
Age	35.2	12.5	18	65	32	25	38	35	0.15	2.1	0.98	Normal
Gender	1.2	0.4	1	2	1	1	1	1	0.05	0.2	0.95	Normal
Marital Status	2.1	0.8	1	3	2	1	3	2	0.12	1.8	0.97	Normal
Education	15.8	2.5	10	20	16	15	17	16	0.08	0.5	0.99	Normal
Income	1200	300	500	2000	1100	800	1400	1000	0.25	1.5	0.96	Normal
Occupation	1.5	0.5	1	3	2	1	3	2	0.10	1.2	0.98	Normal
Health Status	2.5	0.5	1	3	2	1	3	2	0.05	0.2	0.95	Normal
Stress Level	3.2	1.0	1	5	3	2	4	3	0.18	2.5	0.97	Normal
Life Satisfaction	4.1	0.8	3	5	4	3	5	4	0.02	0.1	0.99	Normal
Resilience	3.8	0.9	2	5	4	3	5	4	0.05	0.3	0.98	Normal
Optimism	4.5	0.7	3	5	4	3	5	4	0.01	0.05	0.99	Normal
Emotional Stability	3.5	0.6	2	4	3	2	4	3	0.08	0.4	0.98	Normal
Self-Esteem	4.2	0.5	3	5	4	3	5	4	0.02	0.1	0.99	Normal
Life Satisfaction	4.1	0.8	3	5	4	3	5	4	0.02	0.1	0.99	Normal
Resilience	3.8	0.9	2	5	4	3	5	4	0.05	0.3	0.98	Normal
Optimism	4.5	0.7	3	5	4	3	5	4	0.01	0.05	0.99	Normal
Emotional Stability	3.5	0.6	2	4	3	2	4	3	0.08	0.4	0.98	Normal
Self-Esteem	4.2	0.5	3	5	4	3	5	4	0.02	0.1	0.99	Normal

SEQ ID NO. 100

Met Gln Tyr Phe Glu Trp Tyr Val Pro Asn Asp Gly Glu His Trp Asn Arg Leu Arg Asn Asp Ala  
Glu Asn Leu Ala His Lys Gly Ile Thr Ser Val Trp Ile Pro Pro Val Tyr Lys Gly Thr Ser Gln Asn  
Asp Val Gly Tyr Gly Val Tyr Asp Val Tyr Asp Leu Gly Glu Phe Asn Gln Lys Gly Thr Ile Arg Thr  
Lys Tyr Gly Thr Lys Ala Gln Leu Lys Ser Ala Ile Glu Ala Leu His Asn Gln Asn Ile Asp Val Tyr  
Gly Asp Val Val Met Asn His Lys Gly Gly Ala Asp Tyr Thr Glu Val Val Thr Ala Val Glu Val Asp  
Arg Asn Asn Arg Asn Ile Glu Thr Ser Ser Asp Tyr Gln Ile Asp Ala Trp Thr Gly Phe Asp Phe Pro  
Gly Arg Arg Asp Ser Tyr Ser Asn Phe Lys Trp Arg Trp Phe His Phe Asp Gly Thr Asp Trp Asp Glu  
Gly Arg Lys Leu Asn Arg Ile Tyr Lys Phe Lys Gly Val Gly Lys Ala Trp Asp Trp Glu Val Ser Ser  
Glu Asn Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Phe Asp His Pro Glu Val Ala Asn  
Glu Met Lys Asn Trp Gly Thr Trp Tyr Ala Asp Glu Leu Asn Leu Asp Gly Phe Arg Leu Asp Ala  
Val Lys His Ile Asp His Glu Tyr Leu Arg Asp Trp Val Asn His Val Arg Lys Gln Thr Gly Lys Glu  
Met Phe Thr Val Ala Glu Tyr Trp Gln Asn Asp Ile Arg Thr Leu Asn Asn Tyr Leu Gly Lys Val Asn  
Tyr Asn Gln Ser Val Phe Asp Ala Pro Leu His Tyr Asn Phe His Tyr Ala Ser Thr Gly Asn Gly Asn  
Tyr Asp Met Arg Asn Ile Leu Lys Gly Thr Val Val Glu Ser His Pro Thr Leu Ala Val Thr Leu Val  
Glu Asn His Asp Ser Gln Pro Gly Gln Ser Leu Glu Ser Val Val Ser Pro Trp Phe Lys Pro Leu Ala  
Tyr Ala Phe Ile Leu Thr Arg Ala Glu Gly Tyr Pro Ser Val Phe Tyr Gly Asp Tyr Tyr Gly Thr Asn  
Gly Asn Ser Ser Tyr Glu Ile Pro Thr Leu Lys Asp Lys Ile Asp Pro Ile Leu Thr Ala Arg Lys Asn  
Phe Ala Tyr Gly Thr Gln His Asp Tyr Leu Asp His Pro Asp Val Ile Gly Trp Thr Arg Glu Gly Asp  
Ser Ile His Ala Asn Ser Gly Leu Ala Thr Leu Ile Ser Asp Gly Pro Gly Gly Ser Lys Trp Met Asn  
Val Gly Lys Asn Asn Ala Gly Glu Ile Trp Tyr Asp Ile Thr Gly Asn Gln Thr Asn Thr Val Thr Ile  
Asn Lys Asp Gly Trp Gly Gln Phe His Val Asn Gly Gly Ser Val Ser Ile Tyr Val Gln Lys

SEQ ID NO: 107

atgcaaacgattgcaaaaaaaggggatgaaacgatgaaagggaaaaaatggacagctttagctctaacactgccgctggctgctagcttatca  
acaggcggttcacgccgaaaccgtacataaaggtaaaatctccaacagcagataaaaacgggtgatatttatgagggtgatgtaaactcttttacgatg  
caaataaagatggacatgggtgatttaaaagggtcttacacaaaagtggattattaaatgatggcaattctcatacaaagaatgatctcaagtaa  
gggatttgatgatgccgggtcaacccttctcccagctatcataaatatgatgtaacggactattataattgtatccgcagatggaaatctgcaag  
atttgc aaactgatgaaagaagcagataaacgagatgtaaagtcattatggacctcgttgtgaatcatacgagcagtgaaacaccttggttc  
aagctgcattaaaagataaaaacagcaagtacagagattactatattctgggctgataaaaataccgacttgaatgaaaaaggatcttggggaca  
gcaagtatggcataaagcccaaacggagagtattttacggaacgttttgggaaggaatgccggactaaattacgataatcctgaagtaagaa  
aagaaatgattaacgtaggaaagtgttgctaaagcaaggagttgacgggttcgctatgatgctgcgcttcatttttaaggccaaacacctg

Figure 16 (cont.)

aaggcgctaagaaaaatctctgtgtggaatgaatttagagatgcaatgaaaaaggaaaaccctaacgtatatctaacgggtgaagtatggga  
tcaaccggaagtagtagctcttactatcaatcgcttgattctttattaaactttgatttagcaggaaagattgtaactctgtaaaatcaggaaatgat  
caaggaatcgcgactgcagcagcggcaacggatgaactgttcaaatcatacaatccaaataaaattgacgggtattttctaaccaaccatgacca  
aatcgctcatgagttagtaagcggcgatgtgaataaagcaaaagtcagctgcctctatcttactacgcttctggaacccgtatattttac  
ggtgaagaaatcgcatgaccgggtgaaaagcctgatgagttatccgtgaaccgttccgctggtacgaaggaaacggacttggacaaaccag  
ctgggaaacacctgtatacaaaaaggcggcaacggcgtgtctgtagaagcacaacaaaacaaaaggactctttgttaaatcattaccgtgaa  
atgattcgctgctgcagcagcagcaagagttgtaaaaggaacgcttcaatctatttcagtagacagtaagaagtcgttgcctatagccgcac  
gtataaaggcaaatcgattagcgtgtatcataatatttcaaatcaaccggtaaaagtatctgtagcagcaaaaggtaaatgttttggtagtgaaa  
aagggtgctaagaaagtcaaaaatcagcttgtgattccggcgcaatacaacggttttaataaataa

SEQ ID NO: 168

Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr  
Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ser Pro Thr  
Ala Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His  
Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn  
Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val  
Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala  
Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe  
Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp  
Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly  
Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val  
Gly Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly  
Gln Thr Pro Glu Gly Ala Lys Lys Asn Leu Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys  
Glu Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln  
Ser Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Asn Ser Val Lys Ser Gly Asn Asp  
Gln Gly Ile Ala Thr Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp  
Gly Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala  
Lys Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met  
Thr Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln  
Thr Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln  
Lys Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys  
Gly Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Lys  
Ser Ile Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile  
Phe Gly Ser Glu Lys Gly Ala Lys Lys Val Lys Asn Gln Leu Val Ile Pro Ala Asn Thr Thr Val Leu  
Ile Lys

SEQ ID NO: 169

atgaaaacattcaaattaaaacgcactttttaccgctaaccttgctgctcagtgctcctgcttggctgggcaaaatggcaccatgatgcagtatttt  
cattggtatgtacctaatgatggcgcattatggacgcaggttgaaagcaatgctccagcactcgctgaaacgggtttacagcgctctggctacc  
gccagcttacaaaggcgcggcgagcagtaatgacgtcggttatggcgtctatgatattgacgatttaggtgagttgatcaaaaaggctcagtac  
gaaccaatacggcaccaaggctcagtagatctctgcaatcaatgccgcgcacaacaacaatatccaaatctacggcgatgtgtgtttaaccac  
cgaggttggtgctgatgggaagtcgtgggtcgataccaagcgcgttgattgggacaaccgtaacattgaactgggcgacaatggattgaagct  
tgggttgagtttaattttcctggccgcaacgacaaatactcaacttccattggacttggtatcactttgacgggtgtgactgggatgatgccggcaa  
agaaaaagcgatctttaaattcaaaggcgaaggaaaagcatgggattgggaagtcagctctgaaaaaggcaattacgactacctaattgtacgc  
cgatttagacatggatccaagaagttaacaagagctgaagattggggtgagtggtacatcaacatgaccggcggttgatggcttttagaatg  
gatgccgtgaagcacattaaatatcagtagtacaagagtggttgatgattacgttggaacacaggcaagagctttaccggttggtgagttat  
ggaaftacgacgtaaatcaactgcataactttattactaagacctctggcagtagtgcgttgctgatgcccgcttcacatgaacttcacaacgcg  
tcaaaatctggcggcaattacgatatgcgcaaatcatgaatggcacgttgatgaaggacaaccagtcgaagctgtgactctcgtagaaacc  
acgatacacagccattgcaggcgtagtcgacagtggttggtggttaagcctcttgcttacgattcattttatgcgtgaagaaggttatcc  
atcagtttctacgcagattactacggcgcgagtagcagcgacaaaaggctacaacatcaatatggccaaagttccttacattgaagaactgttaa

Figure 16 (cont.)

cactgcgtaaaagagtatgcgtatggcaaacagaattcttatctcgaccactgggatgtgattggctggacccgagagggcgatgctgaacatcc  
aaactcaatggcggtgatcatgagtatggaccaggtggcaaaaaatggatgtataccggtaagccaagcacgcgctatgtcgacaagctgg  
gtatccgaactgaagaagtttggaccgataccaatggctgggcagaatttcctgtcaatggtggtcagtcctcggttgggtgggcgttaagtaa

SEQ ID NO: 170

Met Lys Thr Phe Lys Leu Lys Arg Thr Phe Leu Pro Leu Thr Leu Leu Ser Ala Pro Ala Phe Ala  
Gly Gln Asn Gly Thr Met Met Gln Tyr Phe His Trp Tyr Val Pro Asn Asp Gly Ala Leu Trp Thr Gln  
Val Glu Ser Asn Ala Pro Ala Leu Ala Glu Asn Gly Phe Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys  
Gly Ala Gly Gly Ser Asn Asp Val Gly Tyr Gly Val Tyr Asp Met Tyr Asp Leu Gly Glu Phe Asp  
Gln Lys Gly Ser Val Arg Thr Lys Tyr Gly Thr Lys Ala Gln Tyr Ile Ser Ala Ile Asn Ala Ala His  
Asn Asn Asn Ile Gln Ile Tyr Gly Asp Val Val Phe Asn His Arg Gly Gly Ala Asp Gly Lys Ser Trp  
Val Asp Thr Lys Arg Val Asp Trp Asp Asn Arg Asn Ile Glu Leu Gly Asp Lys Trp Ile Glu Ala Trp  
Val Glu Phe Asn Phe Pro Gly Arg Asn Asp Lys Tyr Ser Asn Phe His Trp Thr Trp Tyr His Phe Asp  
Gly Val Asp Trp Asp Asp Ala Gly Lys Glu Lys Ala Ile Phe Lys Phe Lys Gly Glu Gly Lys Ala Trp  
Asp Trp Glu Val Ser Ser Glu Lys Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp  
His Gln Glu Val Lys Gln Glu Leu Lys Asp Trp Gly Glu Trp Tyr Ile Asn Met Thr Gly Val Asp Gly  
Phe Arg Met Asp Ala Val Lys His Ile Lys Tyr Gln Tyr Leu Gln Glu Trp Ile Asp His Leu Arg Trp  
Lys Thr Gly Lys Glu Leu Phe Thr Val Gly Glu Tyr Trp Asn Tyr Asp Val Asn Gln Leu His Asn  
Phe Ile Thr Lys Thr Ser Gly Ser Met Ser Leu Phe Asp Ala Pro Leu His Met Asn Phe Tyr Asn Ala  
Ser Lys Ser Gly Gly Asn Tyr Asp Met Arg Gln Ile Met Asn Gly Thr Leu Met Lys Asp Asn Pro Val  
Lys Ala Val Thr Leu Val Glu Asn His Asp Thr Gln Pro Leu Gln Ala Leu Glu Ser Thr Val Asp Trp  
Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Leu Arg Glu Glu Gly Tyr Pro Ser Val Phe Tyr Ala  
Asp Tyr Tyr Gly Ala Gln Tyr Ser Asp Lys Gly Tyr Asn Ile Asn Met Ala Lys Val Pro Tyr Ile Glu  
Glu Leu Val Thr Leu Arg Lys Glu Tyr Ala Tyr Gly Lys Gln Asn Ser Tyr Leu Asp His Trp Asp Val  
Ile Gly Trp Thr Arg Glu Gly Asp Ala Glu His Pro Asn Ser Met Ala Val Ile Met Ser Asp Gly Pro  
Gly Gly Lys Lys Trp Met Tyr Thr Gly Lys Pro Ser Thr Arg Tyr Val Asp Lys Leu Gly Ile Arg Thr  
Glu Glu Val Trp Thr Asp Thr Asn Gly Trp Ala Glu Phe Pro Val Asn Gly Gly Ser Val Ser Val Trp  
Val Gly Val Lys

SEQ ID NO: 171

gtgtatgtaaacctttttacgatgcaataaagatggacatggtgatttaaaaggcttacacaaaagttggattatttaaatgatggcaattctcata  
caaagaatgatcttcaagtaaacgggatttggatgatgccggtcaaccctctcccagctatcataaatatgatgaacggactattataattgat  
ccgcagatggaatctgcaagattttcgaactgatgaagaagcagataaacgagatgtaaaagtcattatggacctcgttgtaacatac  
gagcagtgaaacaccttggtttcaagctgcattaaaagataaaaacagcaagtacagagattactatctgggctgataaaaataccgacttga  
atgaaaaaggatcttggggacagcaagatggcataaaagcccaaacggagagtattttacggaacgttttgggaaggaatgccggacttaaa  
ttacgataatcctgaagtaagaaaagaaatgattaacgtaggaagtttggctaaagcaaggagttgacgggtccgtctagatgctgcgttca  
tatttttaaggccaaacacctgaaggcgctaagaaaaatcctgtggtggaatgaatttagagatgcaatgaaaaaggaaaccctaacgtat  
atctaacgggtgaagtatgggatcaaccggaagtagtagctccttactatcaatcgttgattctttatttaactttgatttagcaggaaagattgtaa  
actctgtaaaatcaggaaatgatcaaggaatcgcgactgcagcagcggaacggatgaactgttcaaatcatacaatccaaataaaattgacgg  
tattttcttaaccaacctgaccaaatacgcgtcatgagtgaactaagcggcgatgtgaataaagcaaagtcagctgcctctacttacttacgctt  
cctggcaaccgctatatttattacggtgaagaaatcggcatgaccggtgaaaagcctgatgagttaatccgtgaaccgttccgctggtacgaagg  
aaacggacttggacaaccagctgggaacacctgtatacaacaagcgcggaacggcggtgtctgtagaagcacaaacaaaaaaggac  
tcttgttaaatcattaccgtgaaatgattcgcgtgcgcagcagcacgaagagttagtaaaaggaacgcttcaatctatttcagtagacagtaaag  
aagtcgttgctatagccgcacgtataaaggcaaatcgattagcgtgtatcataatatttcaaatcaaccggtaaaagtatctgtagcagcaaaag  
gtaaattgatttttgtagtgaagaaagggtgtaagaaagtaaaatcagcttgtgattccggcgaatacaacgggtttaataaaaataa

SEQ ID NO: 172

Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His Gly Asp Leu Lys Gly Leu Thr Gln  
Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn Asp Leu Gln Val Asn Gly Ile Trp  
Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val Thr Asp Tyr Tyr Asn Ile Asp Pro

Figure 16 (cont.)

Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala Asp Lys Arg Asp Val Lys Val  
Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe Gln Ala Ala Leu Lys Asp Lys  
Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp Leu Asn Glu Lys Gly Ser Trp  
Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly Thr Phe Trp Glu Gly Met Pro  
Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val Gly Lys Phe Trp Leu Lys  
Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly Gln Thr Pro Glu Gly Ala  
Lys Lys Asn Leu Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu Asn Pro Asn Val Tyr  
Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser Leu Asp Ser Leu Phe  
Asn Phe Asp Leu Ala Gly Lys Ile Val Asn Ser Val Lys Ser Gly Asn Asp Gln Gly Ile Ala Thr Ala  
Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly Ile Phe Leu Thr Asn  
His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala Lys Ser Ala Ala Ser Ile  
Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr Gly Glu Lys Pro  
Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr Ser Trp Glu Thr  
Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln Lys Asp Ser Leu Leu  
Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly Thr Leu Gln Ser  
Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Lys Ser Ile Ser Val Tyr His  
Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile Phe Gly Ser Glu Lys  
Gly Ala Lys Lys Val Lys Asn Gln Leu Val Ile Pro Ala Asn Thr Thr Val Leu Ile Lys

SEQ ID NO: 173

atgcaaacgattgcaaaaaaggggatgaaacgatgaaagggaataatggacagcttagctctaactgccgctggctgctagcttatca  
acaggcggttcacgcagaaactgtacataaaaggtaaaagctccaacagcagataaaaacgggtgttttatgaggtgtatgaaactcttttacgatg  
caaataaagatggacatgggtgatttaaaggctgtacacaaaagttggattatttaaatgacggcaattctcatacaagaatgatctcaagtaaa  
cgggatttggatgatgccggtaaaccttctcctagctatcataaatatgatgtaacggactattataacattgatcctcagtagcgaagctgcaa  
gatttccgcaaaactgatgaaagaagcagataaacgagacgtaaaagttattatggaccttgtgtgaatcatacagcagtgaaacaccttggtt  
caagctgcaactaaaagataaaaacagcaagtacagagattactatatttgggctgataaaaataccgatttgaatgaaaaaggatcttggggaca  
gcaagtagtgataaaagctccaaacggagagatttttacggaacgttctgggaagggaatgcctgacttaattacgataacctgaagtaagaa  
aagaaatgattaacgtcggaaagtttggctaaagcaaggcggtgatggctccgcttagatgctgcccttcatactttaaagggtcaactcctga  
aggcgctaagaaaaatctcctgtgtggtggaatgagtttagagatgcaatgaaaaagaaaaccctaactatatacgggtgaagtagtgat  
cagccggaagtagtagctccttattatcaatcgcttgattccctatttaactttgatttagcaggaaaaattgtcagctctgtaaaagcaggaaatgat  
caaggaatgccactgcagcagcggcaacggatgagctgttcaaatcatacaatccaaataaaattgacggcattttctaaccaacctgacca  
aaaccgcgtcatgagttagcgaagcggagatgtgaataaagcaaaatcagctgcttcttacttacgcttcttgaaatccgtatatttattacg  
gtgaagaaattggcatgaccgggtgaaaagcctgatgaattaatccgtgaaccgttccgctggtacgaaggcaacggaattggacaaactagct  
gggaaacacctgtatataacaaaggcggcaatgggtgtgtctgtagaagcacaacaaacaaaaggattcttgttaaatcattaccgtgaaatg  
attcgcgtgcgcagcagcacgaagagttagtaaaaggaacgcttcagctctatttcagtagacagtaaaagaagttgtcgttatagccgtacgtat  
aaaggcaactccattagtggtgtatcataatatttcaaataacctgtaaaagtagctgtagcggcgaaaggtaaattgattttgctagtgaaaaagg  
tgctaaaaaaggcaaaaatcagcttggtgattccggcgaaatgcgacggttttaataaaataa

SEQ ID NO: 174

Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr  
Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ala Pro Thr  
Ala Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His  
Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn  
Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val  
Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Ser Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala  
Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe  
Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp  
Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly  
Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val  
Gly Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly

Figure 16 (cont.)

Gln Thr Pro Glu Gly Ala Lys Lys Asn Leu Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys  
Glu Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln  
Ser Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Ser Ser Val Lys Ala Gly Asn Asp  
Gln Gly Ile Ala Thr Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp  
Gly Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala  
Lys Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met  
Thr Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Ile Gly Gln  
Thr Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln  
Lys Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys  
Gly Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Asn  
Ser Ile Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile  
Phe Ala Ser Glu Lys Gly Ala Lys Lys Gly Lys Asn Gln Leu Val Ile Pro Ala Asn Ala Thr Val Leu  
Ile Lys

SEQ ID NO: 175

atgaaaaatataatcagcttgtgtgctgccagcgctatcctcacgggtgtcccacgccagttacgccgacgcaattttacacgcgtttaactggcaat  
ataccgatgtaaccgcaatgcaaatcaaatgcccgaatggctttaaaaaagtcctcattcaccgcaatgaaatccagcgccagtgcaatgg  
tgggcccgtatcaaccgcaagacttgctgtcattgattctccgctgggcaacaacaagatttagtcgcgatgatcaatgcgctcaacagcgt  
tggggctgcagctgtatgctgacgtggtgcttaaccatattggctaacgagtcctggaagcgagtgacctgaactaccggggagtgaggtgct  
caacgactatcaatcccgcagtgcttactatcaaaggcaaacacttttcggcaattacaggagaacctttttccgagaatgatttccatccggca  
ggctgtattaccaatggaaatgatcctggccacgtccagttattggcgcttggtgcggcgacaggcgatactgggctaccggatctcgatcctaa  
tcaatgggttgtagtcagcagaagagttactgaacgcactcaaatcaatgggaatcaaagggttcctgatcgtgagtgcaaacatagagtc  
aatatcaaatagaccaagtgtttaccccagacattaccgctggtatgcataatcggagaagtcattaccagtggtgggcaaggtgatagcggct  
atgaggttttcttgccttaccttaataatccgatcacgccgcttatgacttcccgtatttgcacgattcgagccgcgttttcattctctggtgg  
gttaaatcagctacacaatccacaagcctatggccaagcgttacaggactcacgtgcgatcacctttacgattaccacgacattccaaccaatg  
acggtttccgctaccgatcatgatccaaccgatgaacagctcgctatgcctacatcttgggcaaatgaggaacgccactgtctatagt  
gatgacctactgacagcgaagacaagacagtggtcgttggccgatgtgtggcaagatccgaacatgattaacatgcttgccttcacaacg  
cgatgaaggacaagcatgactgtagtggctagcgatcaatgtacctgtctttaagcgcggaagcaaggcgtggttagaatcaataaatg  
tggcgagagtaagtcggtgactgtcgatacttaccgatgagtttaactggtacacccgtaccaagacgtattgagcggcgacatcaccaca  
gtgagttctcgttatcaccaattgttttggcagcgcgagtgcaaggatgtggaactataa

SEQ ID NO: 176

Met Lys Asn Ile Ile Arg Leu Cys Ala Ala Ser Ala Ile Leu Thr Val Ser His Ala Ser Tyr Ala Asp Ala  
Ile Leu His Ala Phe Asn Trp Gln Tyr Thr Asp Val Thr Ala Asn Ala Asn Gln Ile Ala Ala Asn Gly  
Phe Lys Lys Val Leu Ile Ser Pro Ala Met Lys Ser Ser Gly Ser Gln Trp Trp Ala Arg Tyr Gln Pro  
Gln Asp Leu Arg Val Ile Asp Ser Pro Leu Gly Asn Lys Gln Asp Leu Val Ala Met Ile Asn Ala Leu  
Asn Ser Val Gly Val Asp Val Tyr Ala Asp Val Val Leu Asn His Met Ala Asn Glu Ser Trp Lys Arg  
Ser Asp Leu Asn Tyr Pro Gly Ser Glu Val Leu Asn Asp Tyr Gln Ser Arg Ser Ala Tyr Tyr Gln Arg  
Gln Thr Leu Phe Gly Asn Leu Gln Glu Asn Leu Phe Ser Glu Asn Asp Phe His Pro Ala Gly Cys Ile  
Thr Asn Trp Asn Asp Pro Gly His Val Gln Tyr Trp Arg Leu Cys Gly Gly Gln Gly Asp Thr Gly  
Leu Pro Asp Leu Asp Pro Asn Gln Trp Val Val Ser Gln Gln Lys Ser Tyr Leu Asn Ala Leu Lys Ser  
Met Gly Ile Lys Gly Phe Arg Ile Asp Ala Val Lys His Met Ser Gln Tyr Gln Ile Asp Gln Val Phe  
Thr Pro Asp Ile Thr Ala Gly Met His Ile Phe Gly Glu Val Ile Thr Ser Gly Gly Gln Gly Asp Ser Gly  
Tyr Glu Ala Phe Leu Ala Pro Tyr Leu Asn Asn Thr Asp His Ala Ala Tyr Asp Phe Pro Leu Phe Ala  
Ser Ile Arg Ala Ala Phe Ser Phe Ser Gly Gly Leu Asn Gln Leu His Asn Pro Gln Ala Tyr Gly Gln  
Ala Leu Gln Asp Ser Arg Ala Ile Thr Phe Thr Ile Thr His Asp Ile Pro Thr Asn Asp Gly Phe Arg  
Tyr Gln Ile Met Asp Pro Thr Asp Glu Gln Leu Ala Tyr Ala Tyr Ile Leu Gly Lys Asp Gly Gly Thr  
Pro Leu Val Tyr Ser Asp Asp Leu Pro Asp Ser Glu Asp Lys Asp Ser Gly Arg Trp Ala Asp Val Trp  
Gln Asp Pro Asn Met Ile Asn Met Leu Ala Phe His Asn Ala Met Gln Gly Gln Ser Met Thr Val Val  
Ala Ser Asp Gln Cys Thr Leu Leu Phe Lys Arg Gly Lys Gln Gly Val Val Gly Ile Asn Lys Cys Gly

Figure 16 (cont.)

Glu Ser Lys Ser Val Thr Val Asp Thr Tyr Gln His Glu Phe Asn Trp Tyr Thr Pro Tyr Gln Asp Val  
Leu Ser Gly Asp Ile Thr Thr Val Ser Ser Arg Tyr His Gln Phe Val Leu Pro Ala Arg Ser Ala Arg  
Met Trp Lys Leu

SEQ ID NO: 177

atgaaaacattcaaatataaacgcactttttaccgctgaccttgcctcagtgctcctgcttggcggcaaaatggcaccatgatgcagtattt  
cattggtacgtacctaatgatggcgccattatggacgcaggtgaaagcaatgctccagtactcgtgaaaacggttttacagcgctctggctacc  
gccccatacaaaaggcgccggcgccagtaaatgacgtcggttatggcgctatgatatgtacgatttaggtgagtttgacaaaaaggctcagta  
cgaaccaaatacggcaccaggctcagtaacatctctgcaatcaatccgcgcacacaacaataatccaaatttacggcgacgttggtttaacca  
ccgaggtggcgctgatgggaagtcgtgggtcgataccaagcgcggttgattgggacaaccgcaatattgaactggcgacaaatggattgaag  
cttggttgagtttaatttctggccgcaacgacaatactcgaaactccattggacttggtatcactttgacggtgttgactgggatgatccggc  
aaagaaaaagcgatctttaattcaaaaggcgaaggaaaagcatgggattgggaagtcagctctgaaaaaggcaattacgactacctaattgtac  
gccgatttagacatggatcaccagaaagttaaacaagagctgaaagattggggtgagtggtacatcaacatgaccggcggtgatggcttagaa  
tggtgcccgtgaagcacattaaatatcagtatctacaagagtggtgatcattacgttgaaaacaggcaagagctttaccggttggtgagta  
ttgaattacgacgtaaatcaactgcacaactttattactaagacctctggcagtatgtcgttttcgatgcgccgttcacatgaattctacaacgc  
gtcaaatctggcgccacttacgatatgcgcaaatcatgaatggcacgttgatgaaggacaaccagtcacaagcagtgactctcgtagaaaac  
cacgatacgcagccattgcaggcggttagagtcgacagtagattgggtggtcaagcctctgcttacgcattcattttattgcgtgaagaagggtatc  
catcggtgttctacgcagattactacggcgcgagtcacgcagcaaaaggttacaacattaatatggccaaagtgccttacattgaagaactgtaa  
cactgcgtaaaagagtatgcgtatggcaaacagaattctatctcgaccattgggatgtgattggctggacccgagagggcgatgctgaacatcc  
aaactcaatggcggtgatcatgagtgatggaccggcgccacaaaatggatgtataccggtaagccaagtacgcgctatgtcgacaagctgg  
gtatccgaactgaagatgtttggaccgatgccaatggctgggcagaattctctcaatggtggttcagtcctcggttggtggcggttaagtaa

SEQ ID NO: 178

Met Lys Thr Phe Lys Leu Lys Arg Thr Phe Leu Pro Leu Thr Leu Leu Ser Ala Pro Ala Phe Ala  
Gly Gln Asn Gly Thr Met Met Gln Tyr Phe His Trp Tyr Val Pro Asn Asp Gly Ala Leu Trp Thr Gln  
Val Glu Ser Asn Ala Pro Val Leu Ala Glu Asn Gly Phe Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys  
Gly Ala Gly Gly Ser Asn Asp Val Gly Tyr Gly Val Tyr Asp Met Tyr Asp Leu Gly Glu Phe Asp  
Gln Lys Gly Ser Val Arg Thr Lys Tyr Gly Thr Lys Ala Gln Tyr Ile Ser Ala Ile Asn Ala Ala His  
Asn Asn Asn Ile Gln Ile Tyr Gly Asp Val Val Phe Asn His Arg Gly Gly Ala Asp Gly Lys Ser Trp  
Val Asp Thr Lys Arg Val Asp Trp Asp Asn Arg Asn Ile Glu Leu Gly Asp Lys Trp Ile Glu Ala Trp  
Val Glu Phe Asn Phe Pro Gly Arg Asn Asp Lys Tyr Ser Asn Phe His Trp Thr Trp Tyr His Phe Asp  
Gly Val Asp Trp Asp Asp Ala Gly Lys Glu Lys Ala Ile Phe Lys Phe Lys Gly Glu Gly Lys Ala Trp  
Asp Trp Glu Val Ser Ser Glu Lys Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp  
His Pro Glu Val Lys Gln Glu Leu Lys Asp Trp Gly Glu Trp Tyr Ile Asn Met Thr Gly Val Asp Gly  
Phe Arg Met Asp Ala Val Lys His Ile Lys Tyr Gln Tyr Leu Gln Glu Trp Ile Asp His Leu Arg Trp  
Lys Thr Gly Lys Glu Leu Phe Thr Val Gly Glu Tyr Trp Asn Tyr Asp Val Asn Gln Leu His Asn  
Phe Ile Thr Lys Thr Ser Gly Ser Met Ser Leu Phe Asp Ala Pro Leu His Met Asn Phe Tyr Asn Ala  
Ser Lys Ser Gly Gly Thr Tyr Asp Met Arg Gln Ile Met Asn Gly Thr Leu Met Lys Asp Asn Pro Val  
Lys Ala Val Thr Leu Val Glu Asn His Asp Thr Gln Pro Leu Gln Ala Leu Glu Ser Thr Val Asp Trp  
Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Leu Arg Glu Glu Gly Tyr Pro Ser Val Phe Tyr Ala  
Asp Tyr Tyr Gly Ala Gln Tyr Ser Asp Lys Gly Tyr Asn Ile Asn Met Ala Lys Val Pro Tyr Ile Glu  
Glu Leu Val Thr Leu Arg Lys Glu Tyr Ala Tyr Gly Lys Gln Asn Ser Tyr Leu Asp His Trp Asp Val  
Ile Gly Trp Thr Arg Glu Gly Asp Ala Glu His Pro Asn Ser Met Ala Val Ile Met Ser Asp Gly Pro  
Gly Gly Thr Lys Trp Met Tyr Thr Gly Lys Pro Ser Thr Arg Tyr Val Asp Lys Leu Gly Ile Arg Thr  
Glu Asp Val Trp Thr Asp Ala Asn Gly Trp Ala Glu Phe Pro Val Asn Gly Gly Ser Val Ser Val Trp  
Val Gly Val Lys

SEQ ID NO: 179

atgaaaacattcaaatataaacgcactttttaccgctaaccttgcctcagtgctcctgcttggcgggcaaaatggcaccatgatgcagtactt  
tcattggtacgtacctaatgatggcgccattatggacgcaggtgaaagcaatgctccagcactcgtgaaaacggttttacagcgctctggctacc

Figure 16 (cont.)

gccagcttacaaggcgcggcgccagtaatgatgtcgggttatggcgctctacgatatgtacgatttaggtgagtttgatcaaaaaggtcagtac  
gaaccaaatacggtagcaaggctcagtagatctctgcaatcaatgtcgcgcacaacaataatccaaatttacggcgacgttggtttaaccatc  
gtgttggtcgtgatgggaagtcgtgggtcgataccaagcgcgttgattgggacaaccgtaacattgaactgggcgacaaatggattgaagctt  
gggttgagtttaatttctagccgcaacgacaaatactcgaacttccattggacttggtatcactttgacgggtgtgactgggatgatccgggcaa  
agaaaaagcgatcttaaatcaaggcgaaggaaaagcatgggattgggaagtcagctctgaaaaaggcaattacgactacctaattgtacgc  
cgatttagacatggatcacccagaagttaaacaagagctgaaagattgggtgagtggtacatcaacatgaccggcggttgatggctttagaatg  
gatccgttaagcacaattaaatacagtagctacagaagtgattgatcattacgttggaacacaggcgaagagctttcaccgttggtgagattg  
gaattacgacgttaatacaactgcataactttattactaagacctctggcagtagtgcgtgttcgatgcgccgttcacatgaacttctacaacgcgt  
caaaatctggcggaattacgatatgcgccaatcatgaatggcacgttgatgaaggacaacccagtcgaagctgtgactctcgtagaaaacca  
cgatacgagccattgcaggcggttagtgacagtgattgggttcaagcctctgttgcattcatctgttgcgtgaagaagggttatcca  
tcggtgttctacgcagattactacggcgcgagtagcagcgacaaagggttacaacattaatggccaaagtccttacattgaagaacttgaaca  
ctgcgtaaagagtagtgcgtatggcacaagaattcttatctgcaccattgggatgtgattggctggactcgagagggcgatgctgaacatccaaa  
ctcaatggcggtgatcatgagtgatggaccggcggaacaaatggatgtataccggaatccaagcacgcgtatgtcgacaagctgggtat  
ccgaactgaagatgttggaccgatgccaatggctggcgagaatttctgtcaatgggtggttcagtcctcggttgggtggcggttaagtaa

SEQ ID NO: 180

Met Lys Thr Phe Lys Leu Lys Arg Thr Phe Leu Pro Leu Thr Leu Leu Ser Ala Pro Ala Phe Ala  
Gly Gln Asn Gly Thr Met Met Gln Tyr Phe His Trp Tyr Val Pro Asn Asp Gly Ala Leu Trp Thr Gln  
Val Glu Ser Asn Ala Pro Ala Leu Ala Glu Asn Gly Phe Thr Ala Leu Trp Leu Pro Pro Ala Tyr Lys  
Gly Ala Gly Gly Ser Asn Asp Val Gly Tyr Gly Val Tyr Asp Met Tyr Asp Leu Gly Glu Phe Asp  
Gln Lys Gly Ser Val Arg Thr Lys Tyr Gly Thr Lys Ala Gln Tyr Ile Ser Ala Ile Asn Ala Ala His  
Asn Asn Asn Ile Gln Ile Tyr Gly Asp Val Val Phe Asn His Arg Gly Gly Ala Asp Gly Lys Ser Trp  
Val Asp Thr Lys Arg Val Asp Trp Asp Asn Arg Asn Ile Glu Leu Gly Asp Lys Trp Ile Glu Ala Trp  
Val Glu Phe Asn Phe Pro Ser Arg Asn Asp Lys Tyr Ser Asn Phe His Trp Thr Trp Tyr His Phe Asp  
Gly Val Asp Trp Asp Asp Ala Gly Lys Glu Lys Ala Ile Phe Lys Phe Lys Gly Glu Gly Lys Ala Trp  
Asp Trp Glu Val Ser Ser Glu Lys Gly Asn Tyr Asp Tyr Leu Met Tyr Ala Asp Leu Asp Met Asp  
His Pro Glu Val Lys Gln Glu Leu Lys Asp Trp Gly Glu Trp Tyr Ile Asn Met Thr Gly Val Asp Gly  
Phe Arg Met Asp Ala Val Lys His Ile Lys Tyr Gln Tyr Leu Gln Glu Trp Ile Asp His Leu Arg Trp  
Lys Thr Gly Lys Glu Leu Phe Thr Val Gly Glu Tyr Trp Asn Tyr Asp Val Asn Gln Leu His Asn  
Phe Ile Thr Lys Thr Ser Gly Ser Met Ser Leu Phe Asp Ala Pro Leu His Met Asn Phe Tyr Asn Ala  
Ser Lys Ser Gly Gly Asn Tyr Asp Met Arg Gln Ile Met Asn Gly Thr Leu Met Lys Asp Asn Pro Val  
Lys Ala Val Thr Leu Val Glu Asn His Asp Thr Gln Pro Leu Gln Ala Leu Glu Ser Thr Val Asp Trp  
Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Leu Arg Glu Glu Gly Tyr Pro Ser Val Phe Tyr Ala  
Asp Tyr Tyr Gly Ala Gln Tyr Ser Asp Lys Gly Tyr Asn Ile Asn Met Ala Lys Val Pro Tyr Ile Glu  
Glu Leu Val Thr Leu Arg Lys Glu Tyr Ala Tyr Gly Lys Gln Asn Ser Tyr Leu Asp His Trp Asp Val  
Ile Gly Trp Thr Arg Glu Gly Asp Ala Glu His Pro Asn Ser Met Ala Val Ile Met Ser Asp Gly Pro  
Gly Gly Thr Lys Trp Met Tyr Thr Gly Asn Pro Ser Thr Arg Tyr Val Asp Lys Leu Gly Ile Arg Thr  
Glu Asp Val Trp Thr Asp Ala Asn Gly Trp Ala Glu Phe Pro Val Asn Gly Gly Ser Val Ser Val Trp  
Val Gly Val Lys

SEQ ID NO: 181

ttgccagaggccttcggcctggccattacgccgtcacatagccggcgggggaggttggtggcggtgtcgcgcgggggcagcctgccgatgc  
cggtcctccactggccggcggttcacatcctcgtccggcggttcgtcgcgggtcatccgaacaagcacaagaaccggagatttgcgatgagccaca  
cctcgtgcgcggcgtattggcggtgatcctgctcgggttccccgccctcgtgaccaggccggcaagagcccgccggcggtgcgtacca  
cggcgggcgacgaatcactcctccagggttccactggaacgtcgtcccggaagcgcccaacgactggtacaacatccttcgccagcaggcct  
cgacgatcccgcgagcggttctcggaatctggatgccgggtgcctggcgtgacttctcagctggaccgacggcggaagtcaggcgg  
cggcgaaggctacttctggcacgacttcaacaagaacggcggtacggcgagcgacgccagctgcgccaggccggcgcgactcgggtg  
cgccgggtgaagtgctctacgatgtggtgcccaatcatgaaccggcggtatccggacaaggagatcaacctgccggcgccaggggc  
ttctggcgcaacgactgcaccgacccggcaactacccaacgactgcgatgacggtagccgttcacggcggaagtcggacctgaaca  
ccggccatccgcagatctacggcatgttcgcgacgagcttgccaacctgcgcagcggttacggcgccggcggttccgcttcgacttcgttc

Figure 16 (cont.)

gcggctatgcgcccgaacgggtcgacagctggatgagcgacagcgccgacagcagtttctgcgttggcgagctgtggaaaagcccgtccga  
gtacccgagctgggactggcgcaacacggcgagctggcagcagatcatcaaggactggtccgaccgggccaagtcccgggtgtcgaactc  
gcgctcaaggagcgcatgcagaacggctcggtcgccgactggaagcatggcctcaatggcaacccggaccgcgctggcgcgaggtggc  
ggtgacctttgtcgacaaccacgacaccggctattcgccgggagcagggcgccagcaccactgggcgctgcaggacgggctgatccg  
ccaggcctacgcctacatcctaccagccccgggacgcccgggtggtgtactggtcgacatgtacgactggggctacggcgacttcattcgcca  
gctgatccaggtgcggcgaccgctggcgtgcgcgccgattcgccgacagcttcacagcggtacagcggcctggtcgctaccgtcagc  
ggcagccatcagaccctgggtggtggcgctcaactccgatctggccaacccggccaggtcgccagcgcgagcttcagcgagggcggtcaac  
gccagcaacggccaggtgcgcgtctggcgagcggtagcgcgatggcgggcaatgacggcgcgagggcggtctggtcaatgtgaa  
cttcgctgcgacaacggcggtgacgcagatggcgacagcgctacgcggtgggcaacgtcagccagctcggaactggagccccggcctc  
cgcggtacggctgaccgacaccagcagctatccgacctggaaggcgagcatcgccctgcctgacggtcagaacgtggaatggaagtgcctc  
atccgtaacgaggcgagcgacgctggtgcgccagtggcaatcgggcggaacaaccaggtccaggccgctgccggcgcgagcacca  
gcggctcgttctga

SEQ ID NO: 182

Met Pro Glu Ala Phe Gly Leu Ala Ile Thr Pro Ser His Ser Arg Arg Gly Arg Leu Val Gly Val Ser  
Arg Gly Gly Ser Leu Pro Met Pro Val Leu His Trp Pro Ala Phe Ile Leu Val Arg Arg Phe Val Ala  
Gly His Pro Asn Lys His Lys Asn Arg Ser Ile Ala Met Ser His Thr Leu Arg Ala Ala Val Leu Ala  
Ala Ile Leu Leu Pro Phe Pro Ala Leu Ala Asp Gln Ala Gly Lys Ser Pro Ala Gly Val Arg Tyr His  
Gly Gly Asp Glu Ile Ile Leu Gln Gly Phe His Trp Asn Val Val Arg Glu Ala Pro Asn Asp Trp Tyr  
Asn Ile Leu Arg Gln Gln Ala Ser Thr Ile Ala Ala Asp Gly Phe Ser Ala Ile Trp Met Pro Val Pro Trp  
Arg Asp Phe Ser Ser Trp Thr Asp Gly Gly Lys Ser Gly Gly Gly Glu Gly Tyr Phe Trp His Asp Phe  
Asn Lys Asn Gly Arg Tyr Gly Ser Asp Ala Gln Leu Arg Gln Ala Ala Gly Ala Leu Gly Gly Ala  
Gly Val Lys Val Leu Tyr Asp Val Val Pro Asn His Met Asn Arg Gly Tyr Pro Asp Lys Glu Ile Asn  
Leu Pro Ala Gly Gln Gly Phe Trp Arg Asn Asp Cys Thr Asp Pro Gly Asn Tyr Pro Asn Asp Cys  
Asp Asp Gly Asp Arg Phe Ile Gly Gly Lys Ser Asp Leu Asn Thr Gly His Pro Gln Ile Tyr Gly Met  
Phe Arg Asp Glu Leu Ala Asn Leu Arg Ser Gly Tyr Gly Ala Gly Gly Phe Arg Phe Asp Phe Val  
Arg Gly Tyr Ala Pro Glu Arg Val Asp Ser Trp Met Ser Asp Ser Ala Asp Ser Ser Phe Cys Val Gly  
Glu Leu Trp Lys Ser Pro Ser Glu Tyr Pro Ser Trp Asp Trp Arg Asn Thr Ala Ser Trp Gln Gln Ile Ile  
Lys Asp Trp Ser Asp Arg Ala Lys Cys Pro Val Phe Asp Phe Ala Leu Lys Glu Arg Met Gln Asn  
Gly Ser Val Ala Asp Trp Lys His Gly Leu Asn Gly Asn Pro Asp Pro Arg Trp Arg Glu Val Ala Val  
Thr Phe Val Asp Asn His Asp Thr Gly Tyr Ser Pro Gly Gln Asn Gly Gly Gln His His Trp Ala Leu  
Gln Asp Gly Leu Ile Arg Gln Ala Tyr Ala Tyr Ile Leu Thr Ser Pro Gly Thr Pro Val Val Tyr Trp Ser  
His Met Tyr Asp Trp Gly Tyr Gly Asp Phe Ile Arg Gln Leu Ile Gln Val Arg Arg Thr Ala Gly Val  
Arg Ala Asp Ser Ala Ile Ser Phe His Ser Gly Tyr Ser Gly Leu Val Ala Thr Val Ser Gly Ser His Gln  
Thr Leu Val Val Ala Leu Asn Ser Asp Leu Ala Asn Pro Gly Gln Val Ala Ser Gly Ser Phe Ser Glu  
Ala Val Asn Ala Ser Asn Gly Gln Val Arg Val Trp Arg Ser Gly Ser Gly Asp Gly Gly Gly Asn Asp  
Gly Gly Glu Gly Gly Leu Val Asn Val Asn Phe Arg Cys Asp Asn Gly Val Thr Gln Met Gly Asp  
Ser Val Tyr Ala Val Gly Asn Val Ser Gln Leu Gly Asn Trp Ser Pro Ala Ser Ala Val Arg Leu Thr  
Asp Thr Ser Ser Tyr Pro Thr Trp Lys Gly Ser Ile Ala Leu Pro Asp Gly Gln Asn Val Glu Trp Lys  
Cys Leu Ile Arg Asn Glu Ala Asp Ala Thr Leu Val Arg Gln Trp Gln Ser Gly Gly Asn Asn Gln Val  
Gln Ala Ala Ala Gly Ala Ser Thr Ser Gly Ser Phe

SEQ ID NO: 183

atgcaaacgattgcaaaaaaggggatgaaacgatgaaagggaataatggacagcttagcttaacactgccgctggctgctagcttatca  
acaggcggttcacgccgaaaccgtacataaagtgtaagtctgaagcaacagataaaacgggtgtctttatgaggtgatgtaaacctcttttacgata  
caaataaagatggacatggtgatttaaaaggctcgacacaaaagggtgattatttaaatgacggcaattctacatacaagaatgatcttcaagtaaa  
cgggatttgatgatgccagtaaccctctcctagctatcataaatatgatgtaacggactattataacattgacactcagtcaggaaatctgcaag  
attttcgcaagctgatgaaagaagcagacaaacgagacgtaaaagtcattatggacctgtgtgtaacatacagcagcgaacaccttggtt  
caagctgcattaaaagataaaaaacgaagtagagattactatatttgggctgataaaaataccgatttgaatgaaaaggatcttgggggca  
gcaagtatggcataaagctcaaacggagagtattttacggaacgtttgggaaggaatgcctgacttaattacgataacctgaagtaagaa

Figure 16 (cont.)

aagaaatgattaacgtcggaaagtttggctaaagcaaggcgtaatggcttccgcttagatgctgcgcttcataattttaaggtcaaacacctga  
aggcgctaagaaaaatctgttggtggaatgagtttagagatgcgatgaaaaagaaaaccctaactatataacgggtgaagtatgggat  
cagcctgaagtggtagctccttactatcaatcgcttgattcttatttaatttgatttagcaggaaaaattgtagctctgtaaaagcaggaaatgatc  
aaggaatcgccactgcagcagcggcaacagatgaactgttcaatcatacaatccaaataaaattgacggcattttcttaaccaaccatgaccaa  
aatcgctcatgagtgagctgagcggcgatgtgaacaaagcaaaatcagctgcttctacttacgcttctggcaacccgtatatttattacg  
gtgaagaaattggcatgaccgggtgaaaagcctgatgagttaatccgtgaaccattccgctggtacgaaggaaacggacttggacaaactagct  
gggaaacacctgtatatacaaaaggcggcaacggcgtgtctgtagaagtacaaacaaagggattcttgttaaatcattatcgtgaaatg  
attcgctgctgtagcagcatgaagagttagtaaaaggaacgcttcaatctatttcagtagacagtaagaagtgggtgcctatagtcgcacgtat  
aaaggcaactcattagcgtgtatcataatatttcaaatcaacctgtaaaagtatctgtagcagcgaaaggtaaattgattttgctagtgtgaaaagg  
tgctaaaaaagtcaaaatcagctgttaattccggctaatacaacgggttttaataaaataa

SEQ ID NO: 184

Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr  
Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ser Glu Ala  
Thr Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Thr Asn Lys Asp Gly His  
Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn  
Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val  
Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala  
Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe  
Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp  
Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly  
Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val  
Gly Lys Phe Trp Leu Lys Gln Gly Val Asn Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly  
Gln Thr Pro Glu Gly Ala Lys Lys Asn Ile Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu  
Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser  
Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Ser Ser Val Lys Ala Gly Asn Asp Gln  
Gly Ile Ala Thr Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly  
Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala Lys  
Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr  
Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr  
Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Val Gln Thr Lys Gln Lys  
Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly  
Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Asn Ser Ile  
Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile Phe Ala  
Ser Glu Lys Gly Ala Lys Lys Val Lys Asn Gln Leu Val Ile Pro Ala Asn Thr Thr Val Leu Ile Lys

SEQ ID NO: 185

atgaaactgatgaaagggaataatggacagcttttagctctaactgcccgtggtgctagcttatcaacaggcggttcacgccgaaactgtac  
ataaaggtaaagctccaacagcagataaaaacgggtgtctttatgaggtgtatgtaactcttttacgatgcaataaagatggacatgggtgattta  
aaaggcttacacaaaagctggactatttaaatgacggaaattctcatacaagaatgatcttcaagtaaacgggatttggatgatccagtcac  
ccttctcctagctatcataaatatgatgaacggattattataacattgatccgcagtagcggaaatctgcaagatttgcgaagctgatgaaagaagc  
agacaaacgagacgtaaaagtcattatggaccttgttgtgaatcatagcagcgcgaacaccttggtttcaagctgcgttaaaagataaaaaaca  
gcaagtacagagattactatatttgggctgataaaaataccgacttgaatgaaaaggatcttggggacagcaagtatggcataaagctccaaac  
ggagagtattttacggaacgttttgggaagggaatcctgacttaattacgataaccctgaagtaagaaaagaaatgattaacgtcggaaagttt  
ggctaaagcaaggcgttgatggcttccgcttagatgctgcgcttcataattttaaggtcaaacgcctgaaggcgtaagaaaaatattctgtggt  
ggaatgagtttagagatgcgatgaaaaaagaaaaccctaactatataacgggtgaagtatgggatcagcctgaagtggtagctccttactat  
caatcgcttgattccctatttaactttgatttagcagggaaaattgtcagttctgtaaaagcaggaaatgatcaaggaatcgccactgcagcagcgg  
caacggatgagctgttcaaatcatacaatccaaataaaattgacggcattttcttaaccaaccatgacaaaaccgcgtcatgagtgaactgatcg  
gcgatgtgaacaaagcaaaatcagctgcttcttacttacgcttctggcaacccgtatatttattacgggtgaagaaattggcatgaccgggtga  
aaagcctgatgagttaatccgtgaaccgtccgctggtacgaaggaaacggacttggacaaaccagctgggaaacacctgtatatacaaaagg

Figure 16 (cont.)

cggaacggcgtgtctgtagaagcacaaccaaacaaggattcttgttaaatcattaccgtgaaatgattcgcgtgcgtcagcagcatgaag  
agttagtaaaagggaacgcttcaatctatttagtagacagtaagaagtgtgtgcctatagccgtacgtataaagacaactcgattagcgtgtatcat  
aatatttcaaatcaaccggtaaaagtatctgtagcagcaaaaggtaaatttttctagtgtgaaaagggtgctaaaaagtcagaatcagcttg  
tgattccggctaatacaacggttttaataataa

SEQ ID NO: 186

Met Lys Leu Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr Leu Pro Leu Ala Ala Ser Leu Ser  
Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ala Pro Thr Ala Asp Lys Asn Gly Val Phe Tyr  
Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His Gly Asp Leu Lys Gly Leu Thr  
Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn Asp Leu Gln Val Asn Gly Ile Trp  
Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val Thr Asp Tyr Tyr Asn Ile Asp Pro  
Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala Asp Lys Arg Asp Val Lys Val  
Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe Gln Ala Ala Leu Lys Asp Lys  
Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp Leu Asn Glu Lys Gly Ser Trp  
Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly Thr Phe Trp Glu Gly Met Pro  
Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val Gly Lys Phe Trp Leu Lys  
Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly Gln Thr Pro Glu Gly Ala  
Lys Lys Asn Ile Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu Asn Pro Asn Val Tyr Leu  
Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser Leu Asp Ser Leu Phe Asn  
Phe Asp Leu Ala Gly Lys Ile Val Ser Ser Val Lys Ala Gly Asn Asp Gln Gly Ile Ala Thr Ala Ala  
Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly Ile Phe Leu Thr Asn His  
Asp Gln Asn Arg Val Met Ser Glu Leu Ile Gly Asp Val Asn Lys Ala Lys Ser Ala Ala Ser Ile Leu  
Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr Gly Glu Lys Pro Asp  
Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr Ser Trp Glu Thr Pro  
Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln Lys Asp Ser Leu Leu Asn  
His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly Thr Leu Gln Ser Ile  
Leu Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Asp Asn Ser Ile Ser Val Tyr His  
Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile Phe Ala Ser Glu Lys  
Gly Ala Lys Lys Val Lys Asn Gln Leu Val Ile Pro Ala Asn Thr Thr Val Leu Ile Lys

SEQ ID NO: 187

ttgtatctcatccaggaggggcacatgcgtttccgccattatccaccgcttaccggcctggccgttccggttgagctctgcgtaccgcacag  
agctgcggcataggggagtttgcgacttgcgggttcttgcgaattctgcaaaaaagccggttgcgttgcgtacagcttctccggtcaatgac  
accggcacagaaagtctccatacagcgcgttttgcctttgccttgcacctgtatatacaggcttccgacctgacctgaagcagcgggttc  
gaaaagcagattacagatctgaaaagccggtttgaggacttgccttgcgttgcgtatatacggagctgcgccgtgccaaactggatctctgcgtgc  
agtgtttgataaaaacaaggcaacctcatcgccagtgccgaactggaagcctggatttcagataacccctggatcatcgaatatgcggttttat  
gaaccagaaacaccgcaactttgaagccggctggaacattgggaaaagctgcgcaacccactcataacgaaatacaaaaaacctggcag  
ggtaaaacctggcaggtgacctatcaattcttgcattggctgcagatgcggctggaccagcagtttactgcgccgctacagagtgaacgcc  
ctgggtgtctatcttaaggcgatataacctataatgatgaacgaggattccgcagatgcctgggcgaatccggaattctccgtgacgatctcgg  
gccggaagtccccctgacggtgaaaacccccagggaacaaactggggcttccccattataactgggaaaaccttgcaaatgacgggtacag  
ctggtggaaaaaacgtctgaagcacagcgcacgggtattaccatgcctaccgcattgacctatcttgggttttccggatatgggtataccctat  
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gcgctggcttaccgaacccacttgcctacacgggcagccgaggaagcgaataactgggactatctgggaacacacggctatctgaatcaga  
tcatgaaccgtatcggtgaagaagaactatggctgttcaagcccagatcacctgcgaggcagatatacgaacacaaaacctgccggatgcc  
tgaaagaggttctgttacggcagtggaacacggctgtcgcaggttaccggccgcgacgaaaaggacggacaatctactatccgctgtgg  
cgtttccgtgacagcactgcatggcagacgcttaccgatggcgagaacactccctggaagagctgttcgcccaaaaagcggcgacacatga  
aacctgtggcgagaacaggcgggtggaacttctgggtgagctgacgcgatctacggatgcttgcctgtgctgaagatctgggaagtattccc  
cacagtgtaccggaagtgtttcaaacctttcaattacagcttgcgggttaccgctggggccgcaatgggatgccccggccagcccttca  
cagactggaggagatccgctcatgtcggtagcgaccccatcggtcatgattctctaccctgcgcggatggtgggaacccgaaggcggcgca  
ccgggctttatggacgcatggcctccggaacaggatgcatacgaggagcaggccgcatgagttcgaaggcgctggggaccccgcca

1003137e 02210e

Figure 16 (cont.)

ggcatcctgggtactccgtaaacctctgcgaagcccgtccgcgctctgtgtttcccatccaggatatttggccctgtcttcagactttatgcaat  
gacagcggacgaggaacgcatcaatattccggcagtgatccggattaaactggacataccggttgctgcggcaatcgaggatttatctaaa  
aacagccaactataaccgcaatccagaccgctgcaggaccgccgggcgaggaaggcacaaggagcacagcaatga

SEQ ID NO: 188

Met Tyr Leu Ile Gln Glu Gly His Met Arg Phe Pro Pro Ile Ile His Pro Leu Thr Gly Leu Ala Val Pro  
Val Gly Ala Leu Arg Thr Ala Gln Ser Cys Gly Ile Gly Glu Phe Ala Asp Leu Pro Val Leu Ala Glu  
Phe Cys Lys Lys Ala Gly Phe Asp Leu Val Gln Leu Leu Pro Val Asn Asp Thr Gly Thr Glu Ser Ser  
Pro Tyr Ser Ala Leu Ser Ala Phe Ala Leu His Pro Leu Tyr Ile Arg Leu Ser Asp Leu Pro Glu Ala  
Ala Gly Phe Glu Lys Gln Ile Thr Asp Leu Lys Ser Arg Phe Glu Asp Leu Pro Arg Phe Ser Tyr Thr  
Glu Leu Arg Arg Ala Lys Leu Asp Ile Leu Arg Ala Val Phe Asp Lys Asn Lys Ala Thr Ile Ile Gly  
Ser Ala Glu Leu Glu Ala Trp Ile Ser Asp Asn Pro Trp Ile Ile Glu Tyr Ala Val Phe Met Asn Gln  
Lys His Arg Asn Phe Glu Ala Gly Trp Lys His Trp Glu Lys Leu Arg Asn Pro Thr His Asn Glu Ile  
Gln Lys Thr Trp Gln Gly Lys Thr Trp Gln Ala Asp His Gln Phe Phe Ala Trp Leu Gln Met Arg Leu  
Asp Gln Gln Phe Thr Ala Ala Ala Thr Glu Cys Asn Ala Leu Gly Val Tyr Leu Lys Gly Asp Ile Pro  
Ile Met Met Asn Glu Asp Ser Ala Asp Ala Trp Ala Asn Pro Glu Phe Phe Arg Asp Asp Leu Arg  
Ala Gly Ser Pro Pro Asp Gly Glu Asn Pro Gln Gly Gln Asn Trp Gly Phe Pro Ile Tyr Asn Trp Glu  
Asn Leu Ala Asn Asp Gly Tyr Ser Trp Trp Lys Lys Arg Leu Lys His Ser Ala Arg Tyr Tyr His Ala  
Tyr Arg Ile Asp His Ile Leu Gly Phe Phe Arg Ile Trp Ala Ile Pro Tyr Gly Glu Tyr Ser Gly Tyr Leu  
Gly Trp Pro Leu Pro His Glu Pro Val Ser Ala Ala Glu Leu Ala Glu Arg Gly Phe Ser Lys Asp Arg  
Leu Arg Trp Leu Thr Glu Pro His Leu Pro Thr Arg Ala Ala Glu Glu Ala Asn Asn Trp Asp Tyr Leu  
Gly Thr His Gly Tyr Leu Asn Gln Ile Met Asn Arg Ile Gly Glu Glu Glu Leu Trp Leu Phe Lys Pro  
Glu Ile Thr Cys Glu Ala Asp Ile Arg Asn Thr Asn Leu Pro Asp Ala Leu Lys Glu Val Leu Val Arg  
Gln Trp Lys Asn Arg Leu Leu Gln Val Thr Gly Arg Asp Glu Lys Gly Arg Thr Ile Tyr Tyr Pro Leu  
Trp Arg Phe Arg Asp Ser Thr Ala Trp Gln Thr Leu Thr Asp Gly Glu Lys His Ser Leu Glu Glu Leu  
Phe Ala Gln Lys Ala Ala His Asn Glu Thr Leu Trp Arg Glu Gln Ala Val Glu Leu Leu Gly Glu Leu  
Thr Arg Ser Thr Asp Met Leu Ala Cys Ala Glu Asp Leu Gly Ser Ile Pro His Ser Val Pro Glu Val  
Leu Ser Asn Leu Ser Ile Tyr Ser Leu Arg Val Thr Arg Trp Ala Arg Gln Trp Asp Ala Pro Gly Gln  
Pro Phe His Arg Leu Glu Glu Tyr Pro Leu Met Ser Val Ala Thr Pro Ser Val His Asp Ser Ser Thr  
Leu Arg Gly Trp Trp Glu Thr Glu Gly Gly Asp Arg Ala Phe Met Asp Ala Trp Pro Pro Glu Gln  
Asp Ala Tyr Ala Gly Ala Gly Arg His Glu Phe Glu Gly Ala Trp Gly Pro Arg Gln Ala Ser Trp Val  
Leu Arg Lys Leu Cys Glu Ala Arg Ser Ala Leu Cys Val Phe Pro Ile Gln Asp Ile Leu Ala Leu Ser  
Ser Asp Phe Tyr Ala Met Thr Ala Asp Glu Glu Arg Ile Asn Ile Pro Gly Ser Val Ser Gly Phe Asn  
Trp Thr Tyr Arg Leu Pro Ala Ala Ile Glu Asp Leu Ser Lys Asn Ser Gln Leu Ile Thr Ala Ile Gln Thr  
Ala Leu Gln Asp Arg Arg Ala Arg Lys Ala Gln Gly Ala Gln Gln

SEQ ID NO: 189

atgcaaacgattgcaaaaaaggggatgaaacgatgaaagggaaaaatggacagcttagctctaactgccgctggtgctagcttatca  
acaggcgttacgccgaaaccgtacataaaggtaaattccagctgcagataaaaacggtgtctttatgaggtgatgtaaacctttttacgatg  
caataaagatggacatggtgatttaaaaggcttacacaaaaactggactatttaaatgatggcaattctatacaagaatgatcttcaagtaaa  
cgggatttggatgatgccgatcaacccttctcctagctatcataaatatgatgtaacggactattataacattgattctcagtacggaaatctgcaag  
atttgcgaagctaataaagaagcagataaacgagatgtaaaagtattatggacctcgttgtaatacagagcagtgaaacaccttggttca  
agctcgttaaaagataaaaaacagcaagtacagagattactataatttgggctgataaaaaataccgatttgaatgaaaaaggatcttggggacaac  
aagtatggcacaagctccaaacggagagtattttacggaacgttctgggaaggatgcctgacttaattacgataaacctgaagtaagaaaa  
gaaatgattaacgtcggaaagtgttgctaaagcaaggcgttgacggcttccgcttagatgctgcccttcatactttaaagggtcaaacacctgaa  
ggcgctaagaaaaatattgtgtggtgaatgaatttagagatgcgatgaaaaaagaaaacccgaacgtatatctaacgggcgaagtatgggatc  
agccggaagtgttagctccttattatcagtcgcttgattccctatttaactttgatttagcaggaaaaattgtcagctctgtaaaagcaggaaatgatc  
aaggaatcgctactgcagcagcggcaacagatgaactgttcaaatcataatcaataaaattgacggcattttcttaaccaatcatgacaaaa  
atcgctcatgatgtgagtaagcggagatgtcaataagcaaaagtcagctgcctctatcttactacgcttcttggaatccgtatatttattacggt  
gaagaaatcgcatgaccggtgaaaagcctgatgaattaatccgtgaaccgttccgctggtacgaaggaaacggacttggacaaactagtgg

Variable	Mean	SD	Min	Max	Median	Q1	Q3	Mode	Skewness	Kurtosis	Shapiro-Wilk	Normality
Age	35.2	12.5	18	65	32	28	38	35	0.15	2.1	0.98	Normal
Gender	1.2	0.4	1	2	1	1	1	1	0.05	0.2	0.95	Normal
Education	12.5	2.1	9	16	12	11	13	12	0.12	1.8	0.97	Normal
Income	1500	500	500	3000	1200	800	1800	1000	0.25	3.5	0.92	Normal
Marital Status	1.8	0.6	1	3	2	1	3	2	0.08	0.3	0.96	Normal
Occupation	2.5	1.2	1	5	2	1	3	2	0.18	2.5	0.94	Normal
Health Status	1.5	0.5	1	2	1	1	1	1	0.02	0.1	0.99	Normal
Stress Level	3.2	1.1	1	5	3	2	4	3	0.10	1.9	0.97	Normal
Life Satisfaction	4.5	0.8	3	5	4	4	4	4	0.01	0.05	0.99	Normal
Work-Life Balance	2.8	0.9	1	4	3	2	3	3	0.05	0.2	0.96	Normal
Family Support	3.8	1.0	2	5	4	3	4	4	0.03	0.1	0.98	Normal
Community Involvement	2.2	0.7	1	4	2	1	3	2	0.15	2.0	0.95	Normal
Personal Growth	3.5	0.9	2	5	3	3	3	3	0.02	0.05	0.99	Normal
Overall Well-being	3.0	1.2	1	5	3	2	4	3	0.12	1.8	0.97	Normal

SEQ ID NO: 190

Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr  
Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ser Pro Ala  
Ala Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His  
Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn  
Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Ile Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val  
Thr Asp Tyr Tyr Asn Ile Asp Ser Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala  
Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe  
Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp  
Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly  
Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val  
Gly Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly  
Gln Thr Pro Glu Gly Ala Lys Lys Asn Ile Val Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu  
Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser  
Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Ser Ser Val Lys Ala Gly Asn Asp Gln  
Gly Ile Ala Thr Ala Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly  
Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Asn Lys Ala Lys  
Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr  
Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr  
Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln Lys  
Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly  
Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Asn Ser Ile  
Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile Phe Ala  
Ser Glu Lys Gly Ala Lys Lys Val Lys Asn Gln Leu Val Ile Pro Ala Asn Thr Thr Val Leu Val Lys

atgcaaacgattgcaaaaaaagggggatgaaacgatgaaagggaaaaaatggacagctttagctctaacactgccgctggctgctagcttatca  
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caaataaagatggacatgggtgacttaaaaggcttctacaaaaagttggactattaaatgacggcaattctcatacaaaaaatgatctcaagtaaa  
cgggatttggatgatgccagtcaccccttctcctagctatcataaatatgatgtaacggactattataacattgatccgcagctacggaaatctgcaa  
gattttcgaaagctgatgaaagaagcagacaaacgagacgtaaaagtcattatggaccttgggtgaatcatcagcagtgaaacaccttgggtt  
caagctgcgttaaaaagataaaaacgcaagctacagagattactatatttgggctgataaaaataccgacttgaatgaaaaggatcttggggaca  
acaagtatggcataaaagctccaaacggagagtattttacggaacgttctgggaaggaatgccctgacttaaaftacgataacccctgaagtaagaa  
aagaaatgattaacgtcggaaaagtfttggctaaagcaaggcggtgacgggtccgcttagatgctgcgctcatafttttaagggtcaaacagctga  
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attcgcgtgcgtcagcagcatgaagagttagtaaaaggaacgcttcaatctatttcagtagacagtaaaagaggtgtgcttatagccgtacgtata  
aaggcaactccattagtgtgtatcataatatttcaaatcaaccggtaaaaagtaatctgtagcagcgaagggtaaattgatttttgcgtagtgaaaaagg  
gctaagaaagtcaaaaatcagcttgtgttccggcggaatacaacggttttaataaataa

SEO ID NO: 192

Figure 16 (cont.)

Met Gln Thr Ile Ala Lys Lys Gly Asp Glu Thr Met Lys Gly Lys Lys Trp Thr Ala Leu Ala Leu Thr  
Leu Pro Leu Ala Ala Ser Leu Ser Thr Gly Val His Ala Glu Thr Val His Lys Gly Lys Ser Pro Thr  
Ala Asp Lys Asn Gly Val Phe Tyr Glu Val Tyr Val Asn Ser Phe Tyr Asp Ala Asn Lys Asp Gly His  
Gly Asp Leu Lys Gly Leu Thr Gln Lys Leu Asp Tyr Leu Asn Asp Gly Asn Ser His Thr Lys Asn  
Asp Leu Gln Val Asn Gly Ile Trp Met Met Pro Val Asn Pro Ser Pro Ser Tyr His Lys Tyr Asp Val  
Thr Asp Tyr Tyr Asn Ile Asp Pro Gln Tyr Gly Asn Leu Gln Asp Phe Arg Lys Leu Met Lys Glu Ala  
Asp Lys Arg Asp Val Lys Val Ile Met Asp Leu Val Val Asn His Thr Ser Ser Glu His Pro Trp Phe  
Gln Ala Ala Leu Lys Asp Lys Asn Ser Lys Tyr Arg Asp Tyr Tyr Ile Trp Ala Asp Lys Asn Thr Asp  
Leu Asn Glu Lys Gly Ser Trp Gly Gln Gln Val Trp His Lys Ala Pro Asn Gly Glu Tyr Phe Tyr Gly  
Thr Phe Trp Glu Gly Met Pro Asp Leu Asn Tyr Asp Asn Pro Glu Val Arg Lys Glu Met Ile Asn Val  
Gly Lys Phe Trp Leu Lys Gln Gly Val Asp Gly Phe Arg Leu Asp Ala Ala Leu His Ile Phe Lys Gly  
Gln Thr Ala Glu Gly Ala Lys Lys Asn Ile Leu Trp Trp Asn Glu Phe Arg Asp Ala Met Lys Lys Glu  
Asn Pro Asn Val Tyr Leu Thr Gly Glu Val Trp Asp Gln Pro Glu Val Val Ala Pro Tyr Tyr Gln Ser  
Leu Asp Ser Leu Phe Asn Phe Asp Leu Ala Gly Lys Ile Val Ser Ser Val Lys Ala Gly Asn Asp Gln  
Gly Ile Ala Thr Ala Ala Ala Ala Thr Asp Glu Leu Phe Lys Ser Tyr Asn Pro Asn Lys Ile Asp Gly  
Ile Phe Leu Thr Asn His Asp Gln Asn Arg Val Met Ser Glu Leu Ser Gly Asp Val Ser Lys Ala Lys  
Ser Ala Ala Ser Ile Leu Leu Thr Leu Pro Gly Asn Pro Tyr Ile Tyr Tyr Gly Glu Glu Ile Gly Met Thr  
Gly Glu Lys Pro Asp Glu Leu Ile Arg Glu Pro Phe Arg Trp Tyr Glu Gly Asn Gly Leu Gly Gln Thr  
Ser Trp Glu Thr Pro Val Tyr Asn Lys Gly Gly Asn Gly Val Ser Val Glu Ala Gln Thr Lys Gln Lys  
Asp Ser Leu Leu Asn His Tyr Arg Glu Met Ile Arg Val Arg Gln Gln His Glu Glu Leu Val Lys Gly  
Thr Leu Gln Ser Ile Ser Val Asp Ser Lys Glu Val Val Ala Tyr Ser Arg Thr Tyr Lys Gly Asn Ser Ile  
Ser Val Tyr His Asn Ile Ser Asn Gln Pro Val Lys Val Ser Val Ala Ala Lys Gly Lys Leu Ile Phe Ala  
Ser Glu Lys Gly Ala Lys Lys Val Lys Asn Gln Leu Val Val Pro Ala Asn Thr Thr Val Leu Met Lys

SEQ ID NO: 193

atgaaattcaaaaagagtttatctgccgggctcctttgttcggaggctgagcgggtgacaccatccgtcgtcgggaggtgccacgaaccgc  
atttgccatttattcgaatggagttggccggaattgccaccgaatgcgaacacatttcttgccctaagggggttctctcgggttcaggtgtccgc  
cgcaaaaagcgtcagcaatgctgcctgggtgggcgcgtaccaacctgttagttactctttgaagggcgcagtggaacccgggctcaattgc  
ggatatggtccagcgtgttaaagcgggtgggggtcgatattatctggatgcggtgatcaaccatattggcagcacaagatcgctatttccagaagt  
accttacagcagtaatgattttcacagttgcacgggcgatattcattccaaccgctggctgattcaaaattgcgatctgggtgggctgaacgat  
ctcaaaaccgagtcagaatacgttcggcagaaaattgcagactatatgaacgatgcgctcagctcgtggcggtggcggggtttcggattgatccg  
ccaagcatatccggccggcgacatcgcggcgatcaagagcaagctcaacggcagcccgtatatctatcaggaggttatcggggcgccagg  
ggagccggtacaaaccagcagtagacacgtatattggagacgtgacggaatttaactcggccggaccatcgggcctaaattaaagcaaggtaat  
attaagacctgcaggggattggttcgtggagcggctggctgagcagcgacgatgcggtgacctttgtaccaacctgacgaagaacgcca  
taacctggccaggttctcagccatcaggactttggcaatctgtatttctcggttaacgtgttactctggcgatccttacggctacccaaaagtga  
tgtcggggtactactcagtaattttgatgccggggccaccatcgacaggggtacattctggtaatgcgtgtgcttggatggcggtgattgggtctg  
cgaacacaaatggcggtgttagccaacatgggtggcgtttcgcaaccacacagcagcccagtggcaggtcactgactggtgggacgatggt  
acaatcaggtggcggtttggtcgtggcgggctgggctttgtggtgatcaatcgagatgacaataaaggcatcaatcagagttccagacgggaat  
ggcgctggcgagatattgtgacatattgccggtgatttcgacaccagagcggctattgcagcgctacgacgatcaccgtcgacagtcaggg  
gtatgcacattttactgtcggtatgcatcaggccgctgcgattacattggcgcgaaactcggctccgtgtgccaggactgtggcggcacggcc  
gcagagacaaaagtctgctttgacaatgcacaaaactttagccaaccgtatttgcattactggaatgtcaatgcggatcaggccgtagcgaatgc  
aacctggccggcgctcgcgatgacggctgaaaatggcggttactgtacgattttggtgtcggtctcaattcattcaggtaatttcagcgataa  
cgcgccagccaaaccgctgatctgaccgccagcagtcggacgtgtgttacagaacggaacgtggcgtgacagtgaacttctgtcagagta  
gcaatgtgggcaacgagagttggtatttccgttggaaacctcaaacggttggggcggtgagcgcactcacttatgaggtcgcacaggcctgtaca  
ctacggtgcagagcttaacggggaggagtcgcccgcacgctttaaattgatgatggcaactggagtgcgatccaagtgtgattatcaa  
gtcgggtgattatgccacctacacgatcacgtttgacagccagacgaaggccatcacctgacttcgagtaa

SEQ ID NO: 194

Met Lys Phe Lys Lys Ser Leu Ser Ala Gly Leu Leu Leu Phe Gly Gly Leu Ser Gly Val Thr Pro Ser  
Val Ala Ala Glu Val Pro Arg Thr Ala Phe Val His Leu Phe Glu Trp Ser Trp Pro Asp Ile Ala Thr

Figure 16 (cont.)

Glu Cys Glu Thr Phe Leu Gly Pro Lys Gly Phe Ser Ala Val Gln Val Ser Pro Pro Gln Lys Ser Val  
Ser Asn Ala Ala Trp Trp Ala Arg Tyr Gln Pro Val Ser Tyr Ser Phe Glu Gly Arg Ser Gly Thr Arg  
Ala Gln Phe Ala Asp Met Val Gln Arg Cys Lys Ala Val Gly Val Asp Ile Tyr Leu Asp Ala Val Ile  
Asn His Met Ala Ala Gln Asp Arg Tyr Phe Pro Glu Val Pro Tyr Ser Ser Asn Asp Phe His Ser Cys  
Thr Gly Asp Ile Asp Tyr Ser Asn Arg Trp Ser Ile Gln Asn Cys Asp Leu Val Gly Leu Asn Asp Leu  
Lys Thr Glu Ser Glu Tyr Val Arg Gln Lys Ile Ala Asp Tyr Met Asn Asp Ala Leu Ser Leu Gly Val  
Ala Gly Phe Arg Ile Asp Ala Ala Lys His Ile Pro Ala Gly Asp Ile Ala Ala Ile Lys Ser Lys Leu Asn  
Gly Ser Pro Tyr Ile Tyr Gln Glu Val Ile Gly Ala Ala Gly Glu Pro Val Gln Thr Ser Glu Tyr Thr Tyr  
Ile Gly Asp Val Thr Glu Phe Asn Phe Ala Arg Thr Ile Gly Pro Lys Phe Lys Gln Gly Asn Ile Lys  
Asp Leu Gln Gly Ile Gly Ser Trp Ser Gly Trp Leu Ser Ser Asp Asp Ala Val Thr Phe Val Thr Asn  
His Asp Glu Glu Arg His Asn Pro Gly Gln Val Leu Ser His Gln Asp Phe Gly Asn Leu Tyr Phe Leu  
Gly Asn Val Phe Thr Leu Ala Tyr Pro Tyr Gly Tyr Pro Lys Val Met Ser Gly Tyr Tyr Phe Ser Asn  
Phe Asp Ala Gly Pro Pro Ser Thr Gly Val His Ser Gly Asn Ala Cys Gly Phe Asp Gly Gly Asp Trp  
Val Cys Glu His Lys Trp Arg Gly Val Ala Asn Met Val Ala Phe Arg Asn His Thr Ala Ala Gln Trp  
Gln Val Thr Asp Trp Trp Asp Asp Gly Tyr Asn Gln Val Ala Phe Gly Arg Gly Gly Leu Gly Phe  
Val Val Ile Asn Arg Asp Asp Asn Lys Gly Ile Asn Gln Ser Phe Gln Thr Gly Met Pro Ala Gly Glu  
Tyr Cys Asp Ile Ile Ala Gly Asp Phe Asp Thr Gln Ser Gly His Cys Ser Ala Thr Thr Ile Thr Val  
Asp Ser Gln Gly Tyr Ala His Phe Thr Val Gly Ser His Gln Ala Ala Ala Ile His Ile Gly Ala Lys Leu  
Gly Ser Val Cys Gln Asp Cys Gly Gly Thr Ala Ala Glu Thr Lys Val Cys Phe Asp Asn Ala Gln  
Asn Phe Ser Gln Pro Tyr Leu His Tyr Trp Asn Val Asn Ala Asp Gln Ala Val Ala Asn Ala Thr Trp  
Pro Gly Val Ala Met Thr Ala Glu Asn Gly Gly Tyr Cys Tyr Asp Phe Gly Val Gly Leu Asn Ser Leu  
Gln Val Ile Phe Ser Asp Asn Gly Ala Ser Gln Thr Ala Asp Leu Thr Ala Ser Ser Pro Thr Leu Cys  
Tyr Gln Asn Gly Thr Trp Arg Asp Ser Asp Phe Cys Gln Ser Ser Asn Val Gly Asn Glu Ser Trp Tyr  
Phe Arg Gly Thr Ser Asn Gly Trp Gly Val Ser Ala Leu Thr Tyr Glu Ala Ala Thr Gly Leu Tyr Thr  
Thr Val Gln Ser Phe Asn Gly Glu Glu Ser Pro Ala Arg Phe Lys Ile Asp Asp Gly Asn Trp Ser Glu  
Ser Tyr Pro Ser Ala Asp Tyr Gln Val Gly Asp Tyr Ala Thr Tyr Thr Ile Thr Phe Asp Ser Gln Thr  
Lys Ala Ile Thr Val Thr Ser Gln

SEQ ID NO: 195

atgtgtacagaccgtttctttgatggcgatacatcaaacacgacccttacaaccagaactacgatgctaaaaacgaccggggaacttatcagg  
gcggcgattttaaggaatcacgcaaaaattggattatctcgataagctaggcgtgaacacaatctggatcagcccgatcgtggaaaatacaag  
catgatgtccgttatgacaactctgaaggcattcactatgcttaccacggctactggcaagcaacttcggtgcgttaaacccacacttcggt  
acaatggaagattccatacactgattgacgctgcccatgaaaaaggcatcaagatcatggtgacgtagtattaaaccacactggttatggctta  
aaagatatcaacggagaagtccaatctccagccggttaccgaactgacgcagaacgcagcacatatagcagcctgcttcgccagggttca  
aatgtcggctctgatgaggttggtggcgaattagctggcctacgtgacttaaaaacagaagaccccgagtcgccagacaatcatcgactggc  
aacagactggatcacgaaagctactacagctaaaggaaacacaattgactactccgtgtcgacactgtgaagcacgttgaaagacgcaacat  
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acctgaaacaggtatgatggactcactgcttgactcgactcaaaaggcattgcgcacgatttcgtgaacggcaagcttaaggcagcaaacgat  
gccctgactgcccgcaacggtaaaattgacaacacagctactttaggttcattccttgaagccatgacgaagatggttcctatttaaagaagga  
aatgacaaaggcaagcttaagggtgctgctccctgcaagcaacatcaaaaggccagccggtcatctattatggtgaagacgttggtcaaatg  
gagcaaacactatccgcaatacagataaccgttatgacctggcatgggacaaagtgaacaacacgacgtccttgagcactacactaaggtcct  
gaactcagaagcgctcattcagaagtgttcgctaaagggtgaacgcgcaacaattggcggttctgacgctgataaaattctacttttgcctgtaa  
aatggaacgaagctgctacgtcggttgacgttgctgacacagcaaaagacgtaacactgactgtttctgcaggtgcagtcgtaactgacc  
actatgcagataaaactatactgctcagaagctggagaatcacattgacgatcccgcaaaagctgatggcggtactgtttactaacggttg  
aaggcgagaaatcacagctgctaaagcggaagcggaaggcgacggcacagttgagccagtcctctgcaaccacatccgcattcactacaa  
ccgtacagacaacaactgaaaactacggtgcatggctgtggaacgatgtagcctccctctgccaactggccgactggcgctacaaatgtttg  
aaaaaacagacagctacggtgcatacatcgacgtaccacttaagaggcgctaaagaacatcggttcctctgttatggatgtaacaaaagggtga  
tcagggtaaagacggcgccgacaaagggtttacgatctcatcacctgaaatgaacgaaatttgatcaagcaaggttctgacaaggtgtacactt  
acgagccagttgatctccggcgaaacactgtccggtccactgtacgtgacaacgcagactacgaaaacttcggtatctggaactggggcgga  
tgtaacagcacctccgaaaactggcctacaggcgacgcgaaattcgatggtacagaccgttacggtgcgtatgtcgacattacgctaaaagaa

10081676.000400

ggcgcgaagaacattggaatgattgctcttaacactgcaaattggagagaaagacggcggagataaatccttcaacctctggataaatataatcg  
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cgattctattaggcttcacaaatgactgacggcctaacacctgaattctttaaaggaggtcttgaattaaagattcaactgggtgctgaagttgccatc  
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cttaaatatgggctccgaaagcgagcaaggtaaccgctaactcttggataaaaaataatgccgctgaaaaaatcggcagcgctcgagttaacgaa  
gggtgaaaaaggagcttggtcagctatgggtgctcctggcgacctgaacgtaaccgacttgaagggtattttaccagtatgatgtaacaaatga  
cgggtataactcgccaggtgttagatccttatgaaaatcaatggcagcctttactgtgaatacagaaggcaatgctggtcctgacggggacactg  
ttggcaaggcggcaattcaaaaagcttctcgagagtacttctag

Met Leu Thr Asp Arg Phe Asp Gly Asp Thr Ser Asn Asn Asp Pro Tyr Asn Gln Asn Tyr Asp  
Ala Lys Asn Asn Arg Gly Thr Tyr Gln Gly Gly Asp Phe Lys Gly Ile Thr Gln Lys Leu Asp Tyr Leu  
Asp Lys Leu Gly Val Asn Thr Ile Trp Ile Ser Pro Ile Val Glu Asn Ile Lys His Asp Val Arg Tyr Asp  
Asn Ser Glu Gly His Ser Tyr Tyr Ala Tyr His Gly Tyr Trp Ala Ser Asn Phe Gly Ala Leu Asn Pro  
His Phe Gly Thr Met Glu Asp Phe His Thr Leu Ile Asp Ala Ala His Glu Lys Gly Ile Lys Ile Met  
Val Asp Val Val Leu Asn His Thr Gly Tyr Gly Leu Lys Asp Ile Asn Gly Glu Val Ser Asn Pro Pro  
Ala Gly Tyr Pro Thr Asp Ala Glu Arg Ser Thr Tyr Ser Ser Leu Leu Arg Gln Gly Ser Asn Val Gly  
Ser Asp Glu Val Val Gly Glu Leu Ala Gly Leu Pro Asp Leu Lys Thr Glu Asp Pro Ala Val Arg Gln  
Thr Ile Ile Asp Trp Gln Thr Asp Trp Ile Thr Lys Ala Thr Thr Ala Lys Gly Asn Thr Ile Asp Tyr Phe  
Arg Val Asp Thr Val Lys His Val Glu Asp Ala Thr Trp Met Ala Phe Lys Asn Asp Leu Thr Glu  
Lys Met Pro Thr His Lys Met Ile Gly Glu Ala Trp Gly Ala Ser Ala Asn Asn Gln Leu Gly Tyr Leu  
Glu Thr Gly Met Met Asp Ser Leu Leu Asp Phe Asp Phe Lys Gly Ile Ala His Asp Phe Val Asn Gly  
Lys Leu Lys Ala Ala Asn Asp Ala Leu Thr Ala Arg Asn Gly Lys Ile Asp Asn Thr Ala Thr Leu Gly  
Ser Phe Leu Gly Ser His Asp Glu Asp Gly Phe Leu Phe Lys Glu Gly Asn Asp Lys Gly Lys Leu  
Lys Val Ala Ala Ser Leu Gln Ala Thr Ser Lys Gly Gln Pro Val Ile Tyr Tyr Gly Glu Glu Leu Gly  
Gln Ser Gly Ala Asn Asn Tyr Pro Gln Tyr Asp Asn Arg Tyr Asp Leu Ala Trp Asp Lys Val Glu  
Asn Asn Asp Val Leu Glu His Tyr Thr Lys Val Leu Asn Phe Arg Ser Ala His Ser Glu Val Phe Ala  
Lys Gly Glu Arg Ala Thr Ile Gly Gly Ser Asp Ala Asp Lys Phe Leu Leu Phe Ala Arg Lys Asn Gly  
Asn Glu Ala Ala Tyr Val Gly Leu Asn Val Ala Asp Thr Ala Lys Asp Val Thr Leu Thr Val Ser Ala  
Gly Ala Val Val Thr Asp His Tyr Ala Asp Lys Thr Tyr Thr Ala Ser Glu Ala Gly Glu Ile Thr Leu  
Thr Ile Pro Ala Lys Ala Asp Gly Gly Thr Val Leu Leu Thr Val Glu Gly Gly Glu Ile Thr Ala Ala  
Lys Ala Ala Ser Glu Gly Asp Gly Thr Val Glu Pro Val Pro Ala Asn His Ile Arg Ile His Tyr Asn  
Arg Thr Asp Asn Asn Tyr Glu Asn Tyr Gly Ala Trp Leu Trp Asn Asp Val Ala Ser Pro Ser Ala Asn  
Trp Pro Thr Gly Ala Thr Met Phe Glu Lys Thr Asp Ser Tyr Gly Ala Tyr Ile Asp Val Pro Leu Lys  
Glu Gly Ala Lys Asn Ile Gly Phe Leu Val Met Asp Val Thr Lys Gly Asp Gln Gly Lys Asp Gly Gly  
Asp Lys Gly Phe Thr Ile Ser Ser Pro Glu Met Asn Glu Ile Trp Ile Lys Gln Gly Ser Asp Lys Val  
Tyr Thr Tyr Glu Pro Val Asp Leu Pro Ala Asn Thr Val Arg Val His Tyr Val Arg Asp Asn Ala Asp  
Tyr Glu Asn Phe Gly Ile Trp Asn Trp Gly Asp Val Thr Ala Pro Ser Glu Asn Trp Pro Thr Gly Ala  
Ala Lys Phe Asp Gly Thr Asp Arg Tyr Gly Ala Tyr Val Asp Ile Thr Leu Lys Glu Gly Ala Lys Asn  
Ile Gly Met Ile Ala Leu Asn Thr Ala Asn Gly Glu Lys Asp Gly Gly Asp Lys Ser Phe Asn Leu Leu  
Asp Lys Tyr Asn Arg Ile Trp Ile Lys Gln Gly Asp Asp Asn Val Tyr Val Ser Pro Tyr Trp Glu Gln  
Ala Thr Gly Ile Thr Asn Ala Glu Val Ile Ser Glu Asp Thr Ile Leu Leu Gly Phe Thr Met Thr Asp  
Gly Leu Thr Pro Glu Ser Leu Lys Gly Gly Leu Val Ile Lys Asp Ser Thr Gly Ala Glu Val Ala Ile  
Glu Ser Ala Glu Ile Thr Ser Ala Thr Ser Val Lys Val Lys Ala Thr Phe Asp Leu Glu Lys Leu Pro  
Leu Ser Ile Thr Tyr Ala Gly Arg Thr Val Ser Ala Ser Thr Gly Trp Arg Met Leu Asp Glu Met Tyr  
Ala Tyr Asp Gly Asn Asp Leu Gly Ala Thr Tyr Lys Asp Gly Ala Ala Thr Leu Lys Leu Trp Ala Pro  
Lys Ala Ser Lys Val Thr Ala Asn Phe Phe Asp Lys Asn Asn Ala Ala Glu Lys Ile Gly Ser Val Glu  
Leu Thr Lys Gly Glu Lys Gly Val Trp Ser Ala Met Val Ala Pro Gly Asp Leu Asn Val Thr Asp Leu  
Glu Gly Tyr Phe Tyr Gln Tyr Asp Val Thr Asn Asp Gly Ile Thr Arg Gln Val Leu Asp Pro Tyr Ala

Figure 16 (cont.)

Lys Ser Met Ala Ala Phe Thr Val Asn Thr Glu Gly Asn Ala Gly Pro Asp Gly Asp Thr Val Gly Lys  
Ala Ala Ile Gln Lys Ala Ser Arg Glu Tyr Phe

SEQ ID NO: 197

atgaaaccgtcaaaattcgttttctctctgctgccatcgcttgacgcctctccagtaccgccaatgctgacgccatttgcattgaactggaag  
tactccgacgtcacgcaaacgcctcgcaaatcgcgcggggttataaaaaagtctgattcggcagcactgaaatcgagtggaatgaa  
tggtgggcacgttatcaaccgaagatctgcgctgatcgattccccacttggcaacaaaagtgacttaaatccatgattgatgctctgaaggc  
ggctcgcggttgatgtgtatgccgatgtggtgcttaaccatatggccaatgaaacatggaagcgtgaagacttaaatccctggcagtgaaagtc  
tgcaacaatacgacgtaaacaccagttattatcgggaccaaacgcttttggcaattaacggaaaacctattctctggtttgacttccaccaga  
aggctgtattagcgattggaatgatgccgcaatgttcagtactggcgtcttttggcggtgctggtgaccgagggtgccagacttagatccga  
acaactgggtggtgtcacagcaacgtttgtattgaatgcgctaaaaaggttaggtgtgaaaggctccgcattgatcggttaaacacatgagcc  
aatatcaaatcgaccagattttcactgcagagattaccgccggaatgcacgtgttgggtgaagtatcaccagtggtggcaaggcgactccag  
ctatgagaacttcttagcgcttatctcaacgccaccaaccattcggttacgatttcccactgttgcctctattcgcaacgccttctctacagcg  
tgcatgaacatgcttcatgatccacaagcctatggccaagggtgaaaacgcacgttcaattacctttaccatcacgcacacatcccaacga  
acgacggtttccgttatcaaatcatggatccgaaagatgaagagctggcttacgcttatctcggtaagatggcgccacacctctgatttaca  
gcgacaacttacctgataacgaagatcgatgataatcgccgttgggaagggtgttgaaccgtgacctgatgaagaacatgttgcgttccataac  
caaatgaaggcgcaagagatgacgatgctgtacagcgaccaatgtctactgatgttaagcgcggtaaacaagggtggtcggttaataaat  
gcggtgaagagcgttctacacgttgacacctatcagcatgagttcaactggtatcagccttacacagatacactactggcgtgactgaaacc  
gtgagttcgcgttaccacacctccgaattccagctcgacgcgcgcgatgtacatgctctaa

SEQ ID NO: 198

Met Lys Pro Ser Lys Phe Val Phe Leu Ser Ala Ala Ile Ala Cys Ser Leu Ser Ser Thr Ala Asn Ala  
Asp Ala Ile Leu His Ala Phe Asn Trp Lys Tyr Ser Asp Val Thr Gln Asn Ala Ser Gln Ile Ala Ala  
Ala Gly Tyr Lys Lys Val Leu Ile Ser Pro Ala Leu Lys Ser Ser Gly Asn Glu Trp Trp Ala Arg Tyr  
Gln Pro Gln Asp Leu Arg Val Ile Asp Ser Pro Leu Gly Asn Lys Ser Asp Leu Lys Ser Met Ile Asp  
Ala Leu Lys Ala Val Gly Val Asp Val Tyr Ala Asp Val Val Leu Asn His Met Ala Asn Glu Thr Trp  
Lys Arg Glu Asp Leu Asn Tyr Pro Gly Ser Glu Val Leu Gln Gln Tyr Ala Ala Asn Thr Ser Tyr Tyr  
Ala Asp Gln Thr Leu Phe Gly Asn Leu Thr Glu Asn Leu Phe Ser Gly Phe Asp Phe His Pro Glu  
Gly Cys Ile Ser Asp Trp Asn Asp Ala Gly Asn Val Gln Tyr Trp Arg Leu Cys Gly Gly Ala Gly Asp  
Arg Gly Leu Pro Asp Leu Asp Pro Asn Asn Trp Val Val Ser Gln Gln Arg Leu Tyr Leu Asn Ala  
Leu Lys Gly Leu Gly Val Lys Gly Phe Arg Ile Asp Ala Val Lys His Met Ser Gln Tyr Gln Ile Asp  
Gln Ile Phe Thr Ala Glu Ile Thr Ala Gly Met His Val Phe Gly Glu Val Ile Thr Ser Gly Gly Lys Gly  
Asp Ser Ser Tyr Glu Asn Phe Leu Ala Pro Tyr Leu Asn Ala Thr Asn His Ser Ala Tyr Asp Phe Pro  
Leu Phe Ala Ser Ile Arg Asn Ala Phe Ser Tyr Ser Gly Gly Met Asn Met Leu His Asp Pro Gln Ala  
Tyr Gly Gln Gly Leu Glu Asn Ala Arg Ser Ile Thr Phe Thr Ile Thr His Asp Ile Pro Thr Asn Asp  
Gly Phe Arg Tyr Gln Ile Met Asp Pro Lys Asp Glu Glu Leu Ala Tyr Ala Tyr Ile Leu Gly Lys Asp  
Gly Gly Thr Pro Leu Ile Tyr Ser Asp Asn Leu Pro Asp Asn Glu Asp Arg Asp Asn Arg Arg Trp  
Glu Gly Val Trp Asn Arg Asp Leu Met Lys Asn Met Leu Arg Phe His Asn Gln Met Gln Gly Gln  
Glu Met Thr Met Leu Tyr Ser Asp Gln Cys Leu Leu Met Phe Lys Arg Gly Lys Gln Gly Val Val  
Gly Ile Asn Lys Cys Gly Glu Glu Arg Ser His Thr Val Asp Thr Tyr Gln His Glu Phe Asn Trp Tyr  
Gln Pro Tyr Thr Asp Thr Leu Thr Gly Val Thr Glu Thr Val Ser Ser Arg Tyr His Thr Phe Arg Ile  
Pro Ala Arg Ser Ala Arg Met Tyr Met Leu

SEQ ID NO: 199

gtgagtttgacaaaaaggctcagtagcaaccaatacggcaccaaggctcagtagctctgcaatcaatgccgcgcacaacaacaatatcca  
aatttacggcgatgttgtttaaccaccgaggtggtgctgatgggaagtcgtgggtcgataccaagcgcgttgattgggacaaccgcaatttg  
aactgggcgacaaatggattgaagcttgggttgagtttaatttctggccgcaacgacaaatactcgaacttccattggacttggtatcacttgac  
gggtgttgactgggatgacgccggcaagaaaaagcgatctttaaattcaaaggcgaaggaaaagcatgggattgggaagtcagctctgaaa  
aggcaattacgactacctaa

Figure 16 (cont.)

SEQ ID NO: 200

Val Ser Leu Thr Lys Lys Ala Gln Tyr Glu Pro Asn Thr Ala Pro Arg Leu Ser Thr Ser Leu Gln Ser  
Met Pro Arg Thr Thr Thr Ile Ser Lys Phe Thr Ala Met Leu Cys Leu Thr Thr Glu Val Val Leu Met  
Gly Ser Arg Gly Ser Ile Pro Ser Ala Leu Ile Gly Thr Thr Ala Ile Leu Asn Trp Ala Thr Asn Gly Leu  
Lys Leu Gly Leu Ser Leu Ile Phe Leu Ala Ala Thr Thr Asn Thr Arg Thr Ser Ile Gly Leu Gly Ile  
Thr Leu Thr Val Leu Thr Gly Met Thr Pro Ala Lys Lys Lys Arg Ser Leu Asn Ser Lys Ala Lys Glu  
Lys His Gly Ile Gly Lys Ser Ala Leu Lys Lys Ala Ile Thr Thr Thr

SEQ ID NO: 201

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gtaccagccatcacaaaggcgatctgcaagggatcattgactcgtggattacattcaatcgctggcgctcaatgccattggctaacgccgattt  
ttgaatctattccgggtggaggacaagaccattggcgagacagcgttgatgctacaggctactttgccagtgaactttcaagatagaccctcgt  
ttggcacgttagaacaagcccgtagctggtggaaaaggcacacgcgaaggcttgatgtcttcttgatggagtattggcaccataaaggc  
aatgtggtgccatcaccacaaggtagactgctgctggtgaaaataaccggctcagctaccagagagcctggcgtttacgaagaagtcgcc  
agttactgggtgaaagaggttaagattgatggctggcgctggtatcaagcctatcaagtggcgaccgatgcatggaaagcgatccgtcagagc  
gttgatgaagcgctcacagtcctgaacttatgtgaataacaaagggaaccgtccatccttgggttacatgggtgctgaaatttgaataacgaa  
cgttacatcacagaaccgggttacggcaagaggcgatccggcggtgtgctcggttttgatttccgatgcgttccgagtggtcgaaaccttt  
gcggttaacgaaagtggtgctcagccgaaaaggcgcggaatggttgatgacggcatgtcactgcacagtcagatccggatcatgccaagcct  
aatttaatgttgggcaaccatgatgtggtgctgttggggatctgctgcaacgtggcggtattgctcaccagaacaaccgcaatactggcagcg  
tcataaagcggcgatgtcttcttagcagcgataaccggcccaattacctgtattacggtgaagaaattggcgatcaggttgacggcttgcataa  
aaaatcaaagaagattgtgctgtattggtgtgtgatgaccacgtggcgcgaccagtgcggaagattgatggcggtgacggcgctcactgaatg  
cacagcagctgaactcaaagatatgtcttcttattgatgacattacgtcagcaacatctgcgttatcacaagggggaacgtactaatgtgatggc  
gacagagacagtatacgtagaccataaacaggcagacaatgaagccctgtgtacatggtgagtacgactgataacgcggagtcacacctt  
gaaggggcaagcgattggttcacaaggtgtgctgattgattgtaacgaacgagcggtttatgccaataatggggagatgccaatccattaac  
gggctttggcgacgattcctcaagattgacactccgacagcgcggggtgtgatggcgcaatctgctgcctcggtatcgctaggtgaagg  
gatcatggcccaatgtgatacccaaccggtgaaggcaccgggtccggtagcagaacctgtacgtggttggcgatttggccgatgctggttga  
agcaaaagccgcagcgcggtatcaatacaaaaggcaagcacaatggcgaactgtatcaagtgtgtcgaatgaaaagcggcgccctac  
aagatgcaatacggcacgaaagattggagccacagttactgcagacgggatggcattgaagccgggtaccgcaagtcgctcatagcgggt  
ggctacggtaaagacaccgctgacgttccggaatccggtaagtatgtgtggagcttaacattcagtgatcttggcgagccggagcaaatc  
atggtgtctaagtgtagtaa

SEQ ID NO: 202

Met Thr Ala Lys Ala Asp Asp Leu Arg Ile Tyr Gln Ile Met Val Glu Ser Phe Val Asp Gly Asp Lys  
Gln Val Gly His Gly Thr Gly Tyr Gly Thr Ser His His Lys Gly Asp Leu Gln Gly Ile Ile Asp Ser  
Leu Asp Tyr Ile Gln Ser Leu Gly Val Asn Ala Ile Trp Leu Thr Pro Ile Phe Glu Ser Ile Pro Val Glu  
Gly Gln Asp His Trp Ala Asp Arg Leu Asp Ala Thr Gly Tyr Phe Ala Ser Asp Tyr Phe Lys Ile Asp  
Pro Arg Phe Gly Thr Leu Glu Gln Ala Arg Glu Leu Val Glu Lys Ala His Ala Lys Gly Leu Tyr Val  
Phe Phe Asp Gly Val Phe Gly His His Lys Gly Asn Val Val Pro Ser Pro Gln Gly Arg Leu Pro Val  
Gly Glu Asn Asn Pro Val Ser Tyr Pro Glu Ser Leu Ala Phe Tyr Glu Glu Val Ala Ser Tyr Trp Val  
Lys Glu Leu Lys Ile Asp Gly Trp Arg Leu Asp Gln Ala Tyr Gln Val Pro Thr Asp Ala Trp Lys Ala  
Ile Arg Gln Ser Val Asp Glu Ala Ser Gln Ser Val Thr Tyr Val Asn Asn Lys Gly Glu Thr Val His  
Pro Leu Gly Tyr Met Val Ala Glu Ile Trp Asn Asn Glu Arg Tyr Ile Thr Glu Thr Gly Tyr Gly Lys  
Glu Gly Asp Pro Ala Leu Cys Ser Ala Phe Asp Phe Pro Met Arg Phe Arg Val Val Glu Thr Phe Ala  
Val Asn Glu Ser Gly Val Ser Arg Lys Gly Gly Glu Trp Leu Asn Asp Gly Met Ser Leu His Ser Gln  
Tyr Pro Asp His Ala Lys Pro Asn Leu Met Leu Gly Asn His Asp Val Val Arg Phe Gly Asp Leu  
Leu Gln Arg Gly Gly Ile Ala Ser Pro Glu Gln Pro Gln Tyr Trp Gln Arg His Lys Ala Ala Met Ser  
Phe Leu Ala Ala Tyr Thr Gly Pro Ile Thr Leu Tyr Tyr Gly Glu Glu Ile Gly Asp Gln Val Asp Gly  
Phe Ala Lys Lys Ile Lys Glu Asp Cys Ala Val Ile Gly Leu Cys Asp Asp His Val Ala Arg Thr Ser  
Ala Lys Ile Asp Gly Val Thr Ala Ser Leu Asn Ala Gln Gln Ser Glu Leu Lys Val Tyr Val Ser Ser  
Leu Met Thr Leu Arg Gln Gln His Pro Ala Leu Ser Gln Gly Glu Arg Thr Asn Val Met Ala Thr Glu

Figure 16 (cont.)

Thr Val Tyr Val Asp His Lys Gln Ala Asp Asn Glu Ala Leu Leu Tyr Met Val Ser Thr Thr Asp Asn  
Ala Glu Ser Val Thr Leu Lys Gly Lys Ala Ile Gly Ser Gln Gly Val Leu Ile Asp Leu Leu Thr Asn  
Glu Arg Phe Met Pro Asn Asn Gly Glu Tyr Ala Ile Pro Leu Thr Gly Phe Gly Ala Arg Phe Leu Lys  
Ile Asp Thr Pro Thr Ala Ala Gly Val Met Ala Gln Ser Ala Ala Ser Val Ser Leu Val Gly Glu Gly Ile  
Met Ala Gln Cys Asp Thr Pro Thr Val Glu Gly Thr Gly Pro Val Ala Glu Thr Leu Tyr Val Val Gly  
Asp Phe Ala Asp Ala Gly Trp Lys Gln Lys Pro Gln Arg Ala Tyr Gln Tyr Lys Gly Lys His Asn Gly  
Ser Asn Leu Tyr Gln Val Val Val Asp Glu Lys Ala Gly Ala Tyr Lys Met Gln Tyr Ala Thr Lys Asp  
Trp Ser Pro Gln Phe Thr Ala Asp Gly Met Ala Leu Lys Pro Gly Thr Ala Lys Ser Leu Ile Ala Gly  
Gly Tyr Gly Lys Asp Thr Ala Val Thr Leu Pro Glu Ser Gly Lys Tyr Val Trp Ser Leu Thr Phe Ser  
Asp Leu Gly Glu Pro Glu Gln Ile

SEQ ID NO: 203

atgaagatgaagtcccggtggtgttaggtagtcagtgccatggcgttggcctcttcggcagccaatgccggtgcatggtcacctgtt  
ccagtgaagtacaatgacatgccaacgagtgcgaaagggtgctgggtccaaagggtatgaagcagtcagatcacgccgctgtgaa  
cacctgcaaggctcctcctggtgggtggtctatcagcccgtagctacaagaacttcattctctgggcggtaacgagccgaactcaaaagca  
tgatcgcccggttgaaggccggtggtcaagattacgccgatgcggtattcaaccagctggctggtggtacaggcgtcggtacaggtggtgta  
gcagctacaatgccggcagcttcagctatcccccaattggctacaacgatttcatcacgctgggagcctaccaactatgccgaccgcaacaa  
tgtgcaaaacggtgcctgctggggctgcgggatctggataccggctctgcctatgtgcaggatcagctggtacctatatgaagacctgagt  
ggctgggggtgtggcaggttttctgcttgatgcagcaaaagcatatgacggttggcgtatctctggccatgctcagcaaggcgggcaatcctttgt  
ctactccgaggtgattggtgccacgggtgaaccaatccagccggggaatataccggcattggtgccgtgaccgaatttaatacggcaccga  
tctggcctccaactcaaggggcagatcaagaatctcaagagcatgggagagagctgggggtctgcttgcgtcgaacaaggctgaagctttgtg  
gtcaacatgaccgtgagcggggacatggcggtggcggtatgctgacctacaaggatggtgccctctacaatctggccaacatcttcatgctg  
gcctggccctatggcgctatccccaggtgatgtccggctatgattcggcacaataccgatattggtggccgagcgtacccttgttctcc  
ggctctagctggaactcgcaacaccgctggagcaacatcgccaacatggtctcgttcacaatgccgccaaggcagctccatgaccaactg  
gtgggataatgtaataaccagatcgctttggtgcggcgccaaggcctttgtggtgatcaacaatgaatctccactctgagcaagagcctgc  
agacgggtctgccagccggggagtactgcaacattctggcgggtgatgccctgtgcagcggcagcaccatcaagggtggtgaccagcggtat  
ggccaccttcaacgtggcagggatgaaggcggcagcgatccatcaatgccaagcccgatagcaccagcagtggtgagctcaggtcttctct  
ctggctcttctctctgccaccagtaacaagtttgccagcatgaatctgcggggcaccacaatggctgggcccagcaccgccatgacagtggga  
tgccaaccgtgtctggtcggcgatgtcacctttaccggggcgcggtatgccaatggtgcccagcgcttcaagtttgatgtctatggcaactgg  
acagagagctatggcgatacacaagccgatggcattgccgacaaggggagcgccaaggacatctattcaatggtgtgggcaagtatcgtgtc  
tcgtcaaggagagcgacatgagctacacctgaccagctctccagcaatcaggcaccgggtggcgccatcacccccaagacactctccgt  
caagctgggtgactcagtggtgttcgatgctccgggtccaccgatgatgtgggtgtcactggctacagctggtctaccgggtggcagtgccaag  
accgaaactgtgctgtttgatgctctggttaccgaagaccattaccgtgacagtgccgatggcgttgacctccaaggccagtgccaccg  
tcaccgtcaccgatggcagcggtggttataacagcaactttgccagcctgaacttccgtggcactcccaacagttggggcgccgagccatga  
cgctggtggcagacaacacctgggaggaacgggtcaacttcgatggtcagccaatcagcgcttcaagttcgatatcaagggtgactggagc  
cagaactatggtgatgaacaaggatgggggtggccgaacgtaccgggtgccgatatttacaccactgtgaccggtcaatataaggtgcaattta  
acgactccactttgaagtacacctgaccaagctggccgatagcagcgccaccagctatagcgcgaactttgccagcctctacctgcgtggca  
ccccgaacagctggggcaccaccgcatgaagctggtggccaataacagctggcagggcgaggtgaccttaccggcaaggcgatgccca  
ctggtgcccaacgcttcaagttcgacgtcaagggtgactggagccagaactacgggtgacagcaacatggacgggactgccgaacggactgg  
tggcgatatcaccagtgcctggtgggcacctatctggtgaccttaatgacagcacactgaaatacacctgaccgccaataa

SEQ ID NO: 204

Met Lys Met Lys Ser Arg Ala Trp Leu Leu Gly Ser Ala Val Ala Met Ala Leu Ala Ser Ser Ala Ala  
Asn Ala Gly Val Met Val His Leu Phe Gln Trp Lys Tyr Asn Asp Ile Ala Asn Glu Cys Glu Lys Val  
Leu Gly Pro Lys Gly Tyr Glu Ala Val Gln Ile Thr Pro Pro Ala Glu His Leu Gln Gly Ser Ser Trp  
Trp Val Val Tyr Gln Pro Val Ser Tyr Lys Asn Phe Thr Ser Leu Gly Gly Asn Glu Ala Glu Leu Lys  
Ser Met Ile Ala Arg Cys Lys Ala Ala Gly Val Lys Ile Tyr Ala Asp Ala Val Phe Asn Gln Leu Ala  
Gly Gly Ser Gly Val Gly Thr Gly Gly Ser Ser Tyr Asn Ala Gly Ser Phe Ser Tyr Pro Gln Phe Gly  
Tyr Asn Asp Phe His His Ala Gly Ser Leu Thr Asn Tyr Ala Asp Arg Asn Asn Val Gln Asn Gly  
Ala Leu Leu Gly Leu Pro Asp Leu Asp Thr Gly Ser Ala Tyr Val Gln Asp Gln Leu Ala Thr Tyr Met

Figure 16 (cont.)

Lys Thr Leu Ser Gly Trp Gly Val Ala Gly Phe Arg Leu Asp Ala Ala Lys His Met Ser Val Ala Asp  
Leu Ser Ala Ile Val Ser Lys Ala Gly Asn Pro Phe Val Tyr Ser Glu Val Ile Gly Ala Thr Gly Glu Pro  
Ile Gln Pro Gly Glu Tyr Thr Gly Ile Gly Ala Val Thr Glu Phe Lys Tyr Gly Thr Asp Leu Ala Ser  
Asn Phe Lys Gly Gln Ile Lys Asn Leu Lys Ser Met Gly Glu Ser Trp Gly Leu Leu Ala Ser Asn Lys  
Ala Glu Val Phe Val Val Asn His Asp Arg Glu Arg Gly His Gly Gly Gly Gly Met Leu Thr Tyr Lys  
Asp Gly Ala Leu Tyr Asn Leu Ala Asn Ile Phe Met Leu Ala Trp Pro Tyr Gly Ala Tyr Pro Gln Val  
Met Ser Gly Tyr Asp Phe Gly Thr Asn Thr Asp Ile Gly Gly Pro Ser Ala Thr Pro Cys Ser Ser Gly  
Ser Ser Trp Asn Cys Glu His Arg Trp Ser Asn Ile Ala Asn Met Val Ser Phe His Asn Ala Ala Gln  
Gly Thr Ser Met Thr Asn Trp Trp Asp Asn Gly Asn Asn Gln Ile Ala Phe Gly Arg Gly Ala Lys Ala  
Phe Val Val Ile Asn Asn Glu Ser Ser Thr Leu Ser Lys Ser Leu Gln Thr Gly Leu Pro Ala Gly Glu  
Tyr Cys Asn Ile Leu Ala Gly Asp Ala Leu Cys Ser Gly Ser Thr Ile Lys Val Asp Ala Ser Gly Met  
Ala Thr Phe Asn Val Ala Gly Met Lys Ala Ala Ala Ile His Ile Asn Ala Lys Pro Asp Ser Thr Ser  
Ser Gly Ser Ser Gly Ser Ser Ser Gly Ser Ser Ser Ser Ala Thr Ser Asn Lys Phe Ala Ser Met Asn Leu  
Arg Gly Thr Asn Asn Gly Trp Ala Ser Thr Ala Met Thr Val Asp Ala Asn Arg Val Trp Ser Ala Asp  
Val Thr Phe Thr Gly Ala Ala Asp Ala Asn Gly Ala Gln Arg Phe Lys Phe Asp Val Tyr Gly Asn Trp  
Thr Glu Ser Tyr Gly Asp Thr Gln Ala Asp Gly Ile Ala Asp Lys Gly Ser Ala Lys Asp Ile Tyr Phe  
Asn Gly Val Gly Lys Tyr Arg Val Ser Leu Lys Glu Ser Asp Met Ser Tyr Thr Leu Thr Gln Leu Ser  
Ser Asn Gln Ala Pro Val Ala Ala Ile Thr Pro Lys Thr Leu Ser Val Lys Leu Gly Asp Ser Val Val  
Phe Asp Ala Ser Gly Ser Thr Asp Asp Val Gly Val Thr Gly Tyr Ser Trp Ser Thr Gly Gly Ser Ala  
Lys Thr Glu Thr Val Leu Phe Asp Ala Leu Gly Thr Lys Thr Ile Thr Val Thr Val Ala Asp Ala Asp  
Gly Leu Thr Ser Lys Ala Ser Ala Thr Val Thr Val Thr Asp Gly Ser Val Ala Tyr Asn Ser Asn Phe  
Ala Ser Leu Asn Phe Arg Gly Thr Pro Asn Ser Trp Gly Ala Ala Ala Met Thr Leu Val Ala Asp Asn  
Thr Trp Glu Ala Thr Val Asn Phe Asp Gly Gln Ala Asn Gln Arg Phe Lys Phe Asp Ile Lys Gly Asp  
Trp Ser Gln Asn Tyr Gly Asp Ser Asn Lys Asp Gly Val Ala Glu Arg Thr Gly Ala Asp Ile Tyr Thr  
Thr Val Thr Gly Gln Tyr Lys Val Gln Phe Asn Asp Ser Thr Leu Lys Tyr Thr Leu Thr Lys Leu Ala  
Asp Ser Ser Ala Thr Ser Tyr Ser Ala Asn Phe Ala Ser Leu Tyr Leu Arg Gly Thr Pro Asn Ser Trp  
Gly Thr Thr Ala Met Lys Leu Val Ala Asn Asn Ser Trp Gln Ala Glu Val Thr Phe Thr Gly Lys Gly  
Asp Ala Thr Gly Ala Gln Arg Phe Lys Phe Asp Val Lys Gly Asp Trp Ser Gln Asn Tyr Gly Asp Ser  
Asn Met Asp Gly Thr Ala Glu Arg Thr Gly Gly Asp Ile Thr Ser Ala Val Val Gly Thr Tyr Leu Val  
Thr Phe Asn Asp Ser Thr Leu Lys Tyr Thr Leu Thr Ala Lys

SEQ ID NO: 205

atgtaccgcgtaatacctatttttgattatgagtatgattgtagcttgtgagtcctcaagaaaaaacaaccgaaaccgctcaacctcaacaaa  
tgccgaaaaaacctttgtttgggaggtgccaatgtatatgtttgttaactgaccgtttaacaacggtaaccaacaatgacatcaatttaatag  
gactaaagaatcaggaaaactccgcaattttatgggagggcgatatcaaggcgatcacccaaaaataatgaggggtattttagtaaaactaggc  
gttaatgccatctggcttaccctgggtgtgaacaaatacatggcagtggtgatgaaggtaccggcaatacctatgaccttcatggctattgggcca  
aagattggacaaacttagacccaaatttggcacaaaagaagaccttgcgaactggtggcaactgcccatacgaaggcatcaggatacttta  
gatgtgtaataaaccacaccggcccggttaaccgaccaagacccggttggggagaagattgggtacgtacaggcccgagtgatctatga  
taattacccaataaccaccagtgacacgtggttagccaatttacctgatatacttacagaaagtaataaagtgtggccttaccacctttttaga  
taaatggaaagccgaaggcagattagagcaagaactaaagaactgacgattttttccgcacaggccacccacgcgaccccgcttttac  
attattaaatggcttaccgattacatccgagaatttggggtagatgggttaggggtgataccgtaaaacataccgaagaacggttgggcccag  
ttgatgatgaagccgaattgcttttggcgaatataaaaaagccaaccagacaaggtattggacgataatgaattttatggtaggcgaagtgt  
acaactacgggtatttccggcggaaggttctatgattcggcgataaaaaggtggactatttggaccacggattaaaagccatcaatttgaatg  
aaatatgatgccaattttacctacgatacacttttaggaagtacgatacccttttgcataccaaaacttaaaggcagaagtgctcaactaccttca  
tctcacgacgatggaagtcatttgataaaatgcggcaaaaaccatacagatcggtacaaaattactgctcactccggcgcatcccaatttat  
tacggtgacgaaccgccagaagccttaacatagaaggcgacagggagatgctacgcttcgttctgttatgaattgggaagagctcgagaa  
gacctgccaagcaaaaaatacttcagcattggcaaaaactgggcagtttcaggaacaaccaccccgagttggtcgccgaaggcacaacac  
ccttgcaaaaagccgttttacaccttagcagggtttatcaaaaaatggtttattgacaaagttgtgtagcattagatgccctaaaggccaaa  
aacaattaccgttaattggtgttttgatgacggtacaaaactgttagatgcctattcaggcaagaagaacctcagttaaaatggtatcgttcactt  
cttctgaatttgatattgtttgttagaacaataa

Figure 16 (cont.)

SEQ ID NO: 206

Met Tyr Arg Val Ile Pro Ile Ile Leu Ile Met Ser Met Ile Val Ala Cys Glu Ser Pro Lys Lys Lys Thr  
Thr Glu Thr Ala Gln Pro Ser Thr Asn Ala Glu Lys Pro Phe Val Trp Glu Ala Ala Asn Val Tyr Phe  
Leu Leu Thr Asp Arg Phe Asn Asn Gly Asn Pro Asn Asn Asp Ile Asn Phe Asn Arg Thr Lys Glu  
Ser Gly Lys Leu Arg Asn Phe Met Gly Gly Asp Ile Lys Gly Ile Thr Gln Lys Ile Asn Glu Gly Tyr  
Phe Ser Lys Leu Gly Val Asn Ala Ile Trp Leu Thr Pro Val Val Glu Gln Ile His Gly Ser Val Asp  
Glu Gly Thr Gly Asn Thr Tyr Ala Phe His Gly Tyr Trp Ala Lys Asp Trp Thr Asn Leu Asp Pro Asn  
Phe Gly Thr Lys Glu Asp Leu Ala Glu Leu Val Ala Thr Ala His Ala Lys Gly Ile Arg Ile Leu Leu  
Asp Val Val Ile Asn His Thr Gly Pro Val Thr Asp Gln Asp Pro Val Trp Gly Glu Asp Trp Val Arg  
Thr Gly Pro Gln Cys Thr Tyr Asp Asn Tyr Thr Asn Thr Thr Ser Cys Thr Leu Val Ala Asn Leu Pro  
Asp Ile Leu Thr Glu Ser Asn Glu Asn Val Ala Leu Pro Thr Phe Leu Leu Asp Lys Trp Lys Ala Glu  
Gly Arg Leu Glu Gln Glu Leu Lys Glu Leu Asp Asp Phe Phe Ser Arg Thr Gly His Pro Arg Ala Pro  
Arg Phe Tyr Ile Ile Lys Trp Leu Thr Asp Tyr Ile Arg Glu Phe Gly Val Asp Gly Phe Arg Val Asp  
Thr Val Lys His Thr Glu Glu Thr Val Trp Ala Glu Leu Tyr Asp Glu Ala Val Ile Ala Phe Ala Glu  
Tyr Lys Lys Ala Asn Pro Asp Lys Val Leu Asp Asp Asn Glu Phe Tyr Met Val Gly Glu Val Tyr  
Asn Tyr Gly Ile Ser Gly Gly Arg Phe Tyr Asp Phe Gly Asp Lys Lys Val Asp Tyr Phe Asp His Gly  
Phe Lys Ser Leu Ile Asn Phe Glu Met Lys Tyr Asp Ala Asn Phe Thr Tyr Asp Thr Leu Phe Arg Lys  
Tyr Asp Thr Leu Leu His Thr Lys Leu Lys Gly Arg Ser Val Leu Asn Tyr Leu Ser Ser His Asp Asp  
Gly Ser Pro Phe Asp Lys Met Arg Gln Lys Pro Tyr Glu Ser Ala Thr Lys Leu Leu Leu Thr Pro Gly  
Ala Ser Gln Ile Tyr Tyr Gly Asp Glu Thr Ala Arg Ser Leu Asn Ile Glu Gly Ala Gln Gly Asp Ala  
Thr Leu Arg Ser Phe Met Asn Trp Glu Glu Leu Ala Glu Asp Pro Ala Lys Gln Lys Ile Leu Gln His  
Lys Lys Pro Phe Tyr Thr Phe Ser Arg Val Tyr Gln Lys Asn Gly Phe Ile Asp Lys Val Val Val Ala  
Leu Asp Ala Pro Lys Gly Gln Lys Gln Ile Thr Val Asn Gly Val Phe Asp Asp Gly Thr Lys Leu Val  
Asp Ala Tyr Ser Gly Lys Glu Thr Ser Val Lys Asn Gly Ile Val Ser Leu Ser Ser Glu Phe Asp Ile  
Val Leu Leu Glu Gln Lys

SEQ ID NO: 207

ctgtcgactgagcctttctgtttgggctcgcagactgactctcagccccccccgcagtagctccagacggagtagccgtaatagccgttgccgg  
gtcgtgggcaggggcctcgaggtacacccacccgcttgagtcacccactgttcacccagccgaggttgcggtgtactcgtggtatgc  
acgctcccggaacttcggaacgtagaccacctccggctttgcttgaggcgcaggttgatgtatgtatcagtcgccggttgccttcgtagccgt  
ttctcagaatatacagctcgtcgtgtgctagtaaacgacgctcagtcctcctccggccaggttgcatgtatccagatgaggttcttgagcttatcct  
tgttgagccactcctcgtatgcgcggtagaatattgtcggctggccctcgtagggtgaggtgaacgcgtaggctgatactgttccagattatat  
cggtgtcgtgtgttgaacgaaggttacggccttaaacgggtcgcggctgacgactgtgccccgttcttgagggcctcgacgagtgccgggaa  
tgttctgtgtcaagggccgcgtccatctgttagtagagcgggaagtcgaagacctggcgccgctcgagtagggccagttgaggagtgcatc  
aacgttggtgtccagtagtcccaacggccagccgccccaccagttgagccagtccttgacgacccacgctccgtggcccttcacgtatgc  
aaagcggcaggcatcaacgccgatgctccttaggtaggcggcgtagctctcatcgctcgccagagccagtgctggccagctcttctcgtg  
ggctatgtctgggaagcctccaaatgtgcctcgtcacagcacttgacctcgttgggtggaagtcgaggtagtggcagtatattgcccagg  
ccaccttgagaagtcctccaggtgtatgcccccaacgaacgggttccactcgaggtctccgctcgcggtgtgttatgacgatgtccgctatg  
acctttatgccgtaggcatgggctgtttatcatgttcacgagctcctgcttgagccaaagcgcgtctctaccgttcccttctggtcgtactcacc  
gaggtcaaagaagtcgtagggtcgtagcccatcgaataggcgccgcccattgcccgtcgcgggggaatccaaatggcggaattcccg  
cctcgtaccactcgggtatctgtcctgatggtgtccaccagattcctccacctgggacgtccagtagaaggcctgcattataacgccgccc  
cttcagctcggagtacttgccataagtacctctactagtagattaaaa

SEQ ID NO: 208

Leu Ser Thr Glu Pro Phe Val Leu Gly Ser Arg Leu Thr Leu Ser Pro Pro Arg Ser Ser Ser Arg Arg  
Ser Ser Arg Asn Ser Arg Trp Pro Gly Arg Gly Gln Gly Pro Arg Gly Thr Pro Thr Arg Leu Ser Pro  
Pro Thr Cys Pro Pro Ser Arg Arg Gly Cys Arg Cys Thr Arg Gly Cys Thr Leu Pro Arg Thr Ser Glu  
Arg Arg Pro Thr Phe Arg Leu Cys Leu Arg Arg Gly Cys Met Leu Ser Val Pro Ala Cys Phe Arg

Variable	Mean		SD		t		p	
	Control	Case	Control	Case	Control	Case	Control	Case
Age	30.5	30.5	1.5	1.5	0.0	0.0	1.000	1.000
Gender	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Education	12.0	12.0	1.0	1.0	0.0	0.0	1.000	1.000
Occupation	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Marital status	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Religion	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Income	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Health status	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Family size	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Urban/rural	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Employment	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Marital satisfaction	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Family cohesion	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Parenting style	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child development	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child health	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child behavior	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child cognition	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child emotion	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child social skills	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child academic achievement	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child mental health	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child physical health	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000
Child overall well-being	1.0	1.0	0.0	0.0	0.0	0.0	1.000	1.000

SEQ ID NO: 209

SEQ ID NO: 210

SEQ ID NO: 211

GTGTTTCGTTCTGACACAGTTTCGCGTACCTGCATGTATGGTGCGCTGCGTAATGCCTA  
CCAACCCGATCGGGTGTTTACTGGAGTCACGGTGCGGACATGCAACTTAAAAAAGCAT  
GCTCATCGCCAGGCGCTGTTGTTCATCGTGACGCGGTGCCTGTGCCTGAAATCCAGGC  
AGACCCATAAAAAACAACAACAAACCGATAACAAACGACCCAAGCCTTCTAAGAGGAG  
AAAACGGGATGGCTTTTAAACTACGCAAAAAGGCGCTCGTTGGCCTGTTCACGGCCGG

# Journal of Public Health Management and Practice

SEQ ID NO: 212

VFRSDTVSRTC MYGALRNA YQPDRVFTGVTVRTCNLKKH AHRQALLFIVTRCLCLKSRQT  
HKNNNK PITNDPSLLRGENGMAFKLRKKALVGLFTAGAMVYAGAAASGEIILQGFHWHS  
KWGGNNQGW WQVMEGQANTIANAGFTHVWFPPVHNSADAEGYLPRELNNLNSSYGSEA  
QLRSAIQALNNRGVHAIADVMMNHRVGC SGWADFCNPDWPTWYIVANDSWPGGPKSQN  
WDTGETYHAARDLDHANPQVRNDISHYLSRLKDVGFSGWRWDYAKGFWPGYVGEYN  
WNTNPNFCVGEVWDDLDPNPNPNPHRQQQLVDWVDATGGSCHVFDFTTKGLTNYALQHGG  
YWRLQGDNGGPAGGIGWWPQRMVTFVDNHD TGPSNHC GDGQNLWPVPCDKVMEAYA  
YILTHPGVPSVYWTHFFNWN LGSEISQLMQIRKNQGVHSGSDVWIAEARHGLYAA YINGN  
VAMKMGWDN WSPGWGWSLAASGNNWAVWTR